```
25-MAR-2004.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                             29-MAY-2003.
(NIJJ/) NI J.
(GENT/) GENTZ R.
(RUBE/) RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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RESULT 8
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protein - protein search, using sw model
n on: March 17, 2007, 11:54:03 ; Search time 217 Seconds
(without alignments)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ....SAIQVYMLKSLFEDKRKSRT 229
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01-JUL-2003.
(HUWA-) HUWAN GENOME SCI INC.
(HUWA-) HUWAN GENOME SCI INC.
100.0%; Score 1177; DB 7; Length 229;
lery Match
(ery Match 100.0%; Pred. No. 9.6e-115;
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Pred. No. 9.6e-115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB70497 standard; protein; 229.AA.
Angiogenesis protein AAA4 protein sequence (Fig 4).
WO200111086-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 1177; DB 4;
Pred. No. 9.6e-115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MACCOLL
15-FEB-2001.
(EOSB-) EOS BIOTECHNOLOGY INC.
(EOSB-) EOS BIOTECHNOLOGY INC.
(EOSB-) EOS BIOTECHNOLOGY INC. 9.6e-115;
                                                                                                                                                                                                                                                                 Gapop 10.0 , Gapext 0.5
Searched: 2782304 seqs, 489333398 residues
Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Listing first 1500 summaries
A Geneseq 200701:*
                                                                                                                                                                                                            1 MGDKIWLPFPVLLLAALPPV.....
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Human T1-receptor-(T1R-)like ligand II.
US2002142461-A1.
03-0CT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW48335 standard; protein; 229 AA. Human T1 receptor-like ligand II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAE02806 standard; protein; 229 AA.
Human TIR-like ligand II protein.
WO200142506-A1.
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geneseqp2002s:*
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Maximum DB seq length: 200000000
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1177
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Best Local Similarity
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Best Local Similarity
RESULT 2
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Best Local Similarity
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(RUBE/) RUBEN S M.
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Perfect score:
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RESULT 1
ID AAW9
DE HUT
PN W7
PD Y
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Run on:
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Length 229;
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Human PRO733 (UN0411) protein sequence SEQ ID NO:447.

14-SEP-2000.

(GETH ) GENENTECH INC.
                                                                                                                                                        100.0%; Score 1177; DB 7; 100.0%; Pred. No. 9.6e-115;
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(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 100.0%; Pred. No. 9.6e-115;
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ry Match
t Local Similarity 100.0%; Pred. No. 9.6e-115;
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Human secreted protein encoded by DNA clone vq20
WO200055375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Score 1177; DB 8; 100.0%; Pred. No. 9.6e-115;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human p24 vesicle trafficking protein Hp24-2.
WO9836068-A2.
                                                                                                                                                                                                                                                                                                                                                                                                 ADL91839 standard; protein; 229 AA. Human PRO83480 protein SEQ ID NO:60.
WO2004024076-A2.
ADN35148 standard; protein; 229 AA. Human T1 receptor-like ligand II. US2003100048-A1.
                                                                                                                                                                                                                     ADI34089 standard; protein; 229 AA. Human T1R-like ligand II protein. US2003198618-A1.
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Human PRO733 protein sequence.
WO9946281-A2.
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Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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(INCY-) INCYTE PHARM INC.
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99.5%; Score 1171; DB 3; Length 229; 99.6%; Pred. No. 4.1e-114;
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                                                                                                                                                                                       AB017820 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US20030312156-Ai.
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Novel human secreted and transmembrane protein PRO733
US2002192706-A1.
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Novel human secreted and transmembrane protein PRO733
US2003050239-A1.
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Human secreted and transmembrane PRO polypeptide #18.
US2002177553-A1.
                                                                                                                                                                                                                                                                           99.5%; Score 1171; DB 6; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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(GETH ) GENENTECH INC.
99.5%; Score 1171; DB 6;
                                                                                                                                       99.5%; Score 1171; DB 4; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel secreted and transmembrane protein PRO733.
US2003017563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #205.
US200336180-Al.
                                              AAU12376 standard; protein; 229 AA.
Human PRO733 polypeptide sequence.
WO200140466-A2.
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US2003004311-A1.
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US2002169284-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 17
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Best Local Similarity
RESULT 23
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Query Match
Best Local Similarity
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Best Local S
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                                 RESULT 15
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Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003022328-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU67050 standard; protein; 229 AA.
Human secreted/transmembrane, PRO, protein SEQ ID
US2003032155-A1.
                                                                                                    20-FEB-2003.
(GETH ) GENENTECH INC.
Pry Match 99.5%; Score 1171; DB 6;
Pred. No. 4.1e-114;
Best Local Similarity 99.6%; Pred. No. 4.1e-114; RESULT 24
                                                                                                                                                                                                                                                                                    Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                         ABO25045 standard; protein; 229 AA.
Human secreted/transmembrane protein (PRO) #205.
US2003036179-A1.
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                           Human p24 vesicle trafficking protein Hp24-2.
US2002132999-A1.
                                                                                                                                                                                         ABUB0409 standard; protein; 229 AA.
Human secreted/transmembrane protein PRO733.
US2003004102-A1.
                                                                                                                                                                                                                                                                                                                                          ABG72289 standard; protein; 229 AA.
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Human PRO polypeptide #205.
US2003073212-Al.
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02-0AN-2003.
(GETH ) GENENTECH INC.
99.5$;
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ry Match
t Local Similarity 99.6%;
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ry Match 99.5%;
t Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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US2003049816-A1.
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OJ-WAY-2-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
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                                                                                                                                               Novel human secreted and transmembrane protein PRO733.
US2003082711-A1.
01-MAY-2003.
GETH ) GENENTECH INC.
99.5%; Score 1171; DB 6; Lengt
st Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADABSB86 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003082693-A1.
                                                                                                                                                                                                                                                                                                                           99.5%; Score 1171; DB 6; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #205.
US2003068794-Al.
RESULT 33
LD ADB2759 standard; protein; 229 AA.
DE Human PRO polypeptide #205.
PN US2003082704-AL.
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US2003087350-A1.
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US2003082763-A1.
                                                                                                                                                                                                                                                                                                                                                                       ADA47788 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003073215-A1.
                                                                                                                                  protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 229 AA
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US2003082705-A1.
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10-APR-2003.
(GETH ) GENENTECH INC.
99.5%;
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US2003068795-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 37
ID ADACTOR
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                                                                                                Best Local Similarity RESULT 34
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Best Local Similarity
RESULT 42
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Best Local Similarity
                                                                                                                                  ADA86438 standard;
                                                                                        Query Match
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01-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
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ADA87541 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003087345-A1.
                                                                                                                                                                                                                                                                                     AAA31835 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082694-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003082691-A1.
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08-MAY-2003.

(GETH) GENENTECH INC.

99.5%; Score 1171; DB 6;

ery Match

199.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                           Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                  Human PRO polypeptide #205.
US2003087349-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB14898 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB13282 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA94074 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003077722-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB18859 standard; protein; 229 AA
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17-APR-2003.
(GETH ) GENENTECH INC.
99.5%; S
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PA (GETH ) GENENTECH INC.

QUETY MATCh 99.5%;

BEST Local Similarity 99.6%;
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                                                      08-MAY-2003.

1 (GETH) GENENTECH INC.

Query Match

Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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RESULT 51
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Best Local Similarity
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                                                                                                                              RESULT 43
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RESULT

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Query Match ,99.5%; Score 1171; DB 6; Length 229; Best Local Similarity 99.6%; Pred. No. 4.1e-114;
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Human PRO polypeptide SEQ ID NO 410
US2003077714-A1.
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17-APR-2003.
(GETH ) GENENTECH INC.
Match '...tty 99.6%;
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24-APR-2003.
(GETH ) GENENTECH INC.
99.5%;
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24-APR-2003.
(GETH ) GENENTECH INC.
MATCh 17-11 99.5%;
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RESULT 63
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Human PRO polypeptide #205.
US2003096386-A1.
                                                                                                                                                                                                                                                                               01-MAY-2003.
(GETH ) GENENTECH INC.
                 17-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 64
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Best Local Similarity
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Best Local Similarity
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RESULT 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                               ADA80566 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
US2003049817-A1.
US2003073214-A1.
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733. US2003044945-A1.
                                                                                                              ABO19711 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003050240-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA85334 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US200308265-A1.
US2003.
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Novel human secreted and transmembrane protein PRO733
US2003082708-A1.
                                                               Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                                                                                Score 1171; DB 6;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                Score 1171; DB 6;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                ADA12647 standard; protein; 229 AA.
Human secreted/transmembrane polypeptide PRO733.
US2003055216-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB24769 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077713-Al.
                                                                                                                                                                                                                                                                                                                                                                              ADA74536 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003068798-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA82293 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003082701-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA75256 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003073216-A1.
17-APR-2003.
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Human PRO polypeptide #205.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ II.

Lety Match GENENTECH INC.

Best Local Similarity 99.6%; Pre RESULT 56

ID ADA82293 standard; prot PN US200308270**

PN US200308270**

PA OL-MP**
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10-APR-2003.
(GETH ) GENENTECH INC.
99.5%; SC
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                20-MAR-2003.
(GETH ) GENENTECH INC.
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LOST MATCH

Best Local Similarity >
RESULT 57

ID ADA72256 standa

DE Human PRO

PN US20030

PD 17
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Best Local Similarity 1

RESULT 59

ID ADA84782 standar

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PN US20030^*

PA 01-**
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                                               (GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 55
ID ADB24769 standard; pr.
DE Human PRO polypeptide
PN US2003077713-A1.
PD 24-APR-2003.
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                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 54
                                                                           Best Local Similarity
RESULT 52
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RESULT 60
ID ADB30038 standard,
DE Human PRO polvoepti
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                                  06-MAR-2003
                                                                  Query Match
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99.5%; Score 1171; DB 6; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                                 Length 229;
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(GETH) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 6; Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 229;
    DE Human PRO POLYPEPLIGE #205.

PN US2003082761-A1.

PD 01-MAY-2003.

PA (GETH ) GENENTECH INC.

Querry Match

Best Local Similarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human PRO polypeptide SEQ ID NO 410
US2003077715-A1.
                                                                                                                                                                                                                                                                                                  Human PRO polypeptide #205.
17-App.
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Human PRO polypeptide #205.
US2003077721-A1.
                                                                                                                                                      ADA75808 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082703-A1.
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ADB26855 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003092147-Al.
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protein; 229 AA
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Query Match
Best Local Similarity
RESULT 79
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RESULT 88
                                                                                        01-MAY-200.
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Best Local S
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                             99.5%; Score 1171; DB 6; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO19602 standard; protein; 229 AA.
Novel human secreted and transmembrane polypeptide #70.
US2003049633-A1.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA86990 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.01203082709-A1.
01-01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB21788 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733 US2003082765-A1.
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                Human PRO polypeptide #205.
US2003082702-A1.
                                                                    ADA96546 standard; protein; 229 AA. Human PRO polypeptide #205. US2003002690-A1. O1-MAY-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PRO polypeptide #205.
US2003068797-Al.
                                                                                                                                                                                                                                                                                                          ADA95994 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082759-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                          ADB26303 standard; protein; 229 AA. Human PRO polypeptide #205. US2003082760-A1.
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US2003077710-A1.
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01-MAY-2003.
(GETH ) GENENTECH INC.
%**rch 99.5%; Sc
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01-MAY-2003.
(GETH ) GENENTECH INC.
43rCh 99.5%; S/
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01-MAY-2003.
(GETH ) GENENTECH INC.
Watch 99.5%; Sr
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24-APR-2003.
(GETH ) GENENTECH INC.
99.5%; Sr
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
RESULT 77
24-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                    01-MAY-2003.
(GETH ) GENENTECH INC.
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                                         Best Local Similarity RESULT 70
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RESULT 71
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Best Local Similarity
RESULT 73
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Best Local Similarity
RESULT 74
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Best Local Similarity
RESULT 76
                                Query Match
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99.5%; Score 1171; DB 7; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                ADA88093 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082700-A1.
                                                                                                                                                                                          ADA46481 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003054516-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         אראפינום standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003073213-A1.
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Pred. No. 4.1e-114;
                                                                                                                                         Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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US2003022239-A1.
30-OAN-2003.
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                                                                                                                                                                                                                                                                                                                                ADB28511 standard; protein; 229 AA. Human PRO polypeptide #205. US2003082699-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB29063 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003082706-A1.
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Human PRO polypeptide #205.
US2003059909-A1.
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                                                                                                                         (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                            20-MAR-2003
(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                    Best Local Similarity RESULT 80
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Human PRO polypeptide #205.
US2003082762-A1.
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                                                                                        Score 1171; DB 7; Length 229;
Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                          Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA92387 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB38702 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003082766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB30150 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733
US2003087347-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB66622 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003082689-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                      99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                         ADB23665 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077712-A1.
24-APR-2003.
(GETH.) GENENTECH INC.
ADA67031 standard; protein; 229 AA. Human PRO polypeptide #205. US2003068793-A1.
                                                                                                                                       JA DB22892 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003077711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB15450 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003087352-A1.
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
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Human PRO polypeptide #205.
US2003082698-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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RESULT 97
ID ADB90434 standard. ~
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Best Local Similarity
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Best Local Similarity
RESULT 96
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Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                ADB47158 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB77370 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US200308269.A1.
01-MAY-2003.
                                                                                                                           Novel human secreted and transmembrane protein PRO733 US2003082764-A1.
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Pred. No. 4.1e-114;
                                                       Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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US2003077719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB34527 standard; protein; 229 AA. Human PRO polypeptide SEQ ID NO 410. US20030777717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB33975 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410
                                                                                                              protein; 229 AA.
                                                                                                                                                                                                                                                         protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB76669 standard; protein; 229 AA.
Human PRO polypeptide #70.
US2003083248-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 229 AA
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24-APR-2003.
(GETH) GENENTECH INC.
99.5%;
US2003002...
01-MAY-2003.
(GETH ) GENENTECH INC. 99.5%;
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01-MAY-2003.
(GETH ) GENENTECH INC.
""red 99.5%;
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(GETH ) GENENTECH INC.
Query Match 99.5%;
Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
2ry Match 99.5%;
it Local Similarity 99.6%;
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Human PRO polypeptide #205.
US2003082697-A1.
                                                                                                                                                                                                                                                      ADB73953 standard; proteir
Human PRO polypeptide #70.
US2003045462-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 99
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GETH ) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
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Best Local Similarity
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 10-APR-2003
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                                                                           RESULT 116
                                                                                                                                                                                                                                                                                                                                                                             RESULT 118
                                      Query Match . 99.5%; Score 1171; DB 7; Length 229;
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 107
                                                                                                                                                                                                      99.5%; Score 1171; DB 7; Length 229; 99.6%; Pred. No. 4.1e-114;
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PA (GETH) GENENTECH INC.

Query Match 99.5%; Score 1171; DB 7; Length 229;

Best Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 113
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PA (GETH ) GENENTECH INC.

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99.5%; Score 1171; DB 7; Length 229;

Best Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 114

ID ADCEO9043 standard; protein; 229 AA.

DE Human secreted/transmembrane protein, PRO733.
                                                                                                                                                                                                                                                                                                                                                        Score 1171; DB 7; Length 229;
Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                            ADB46578 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082692-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003054405-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC44095 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003054986-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC61855 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003049684-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC66919 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003060406-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC63103 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
                                                                                                                                                                                                                                                        ADB36183 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077720-A1.
24-APR-2003.
(GFTH ) GENENTECH INC.
Duery Match
99.5%; Score 117
est Local Similarity 99.6%; Pred. No.
                                                                                                Human PRO polypeptide SEQ ID NO 410.
US200307718-A1.
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20-MAR-2003.
(GETH ) GENENTECH INC.
Warch 99.5%; Sc
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13-MAR-2003.
(GETH ) GENENTECH INC.
Warch 99.5%; SC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 112
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RESULT 109
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Best Local Similarity
RESULT 110
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Best Local Similarity
RESULT 111
US2003077716-A1.
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Length 229;
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Novel human secreted and transmembrane protein Seg ID410.
US2003087355-A1.
08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO733. US2003092107-A1.
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 Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                         ADC41488 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
17-APR-2003.
10 GETH) GENBNTECH INC.
99.5%; Score 1171; DB
Best Local Similarity 99.6%; Pred. No. 4.1e-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC42112 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2001104998-A1.
GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                   Human secreted/transmembrane protein, PRO733. US2003073131-Al.
                                            ADC68168 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003069178-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein, PRO733. US2003073624-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 229 AA
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15-MAY-2003.
(GETH ) GENENTECH INC.
MATCh '-'larity 99.6%;
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RESULT 121
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PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 99.6%;
                                                                                                    10-APR-2003.
(GETH) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
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Best Local Similarity 99.6%;
RESULT 120
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Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Length 229;

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RESULT 125

RESULT

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Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003087354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC80595 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003092103-A1.
IS-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                           ADC00227 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003087348-A1.
                                                 ADD01235 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003092104-A1.
Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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99.5%; Score 1171;
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US2003194771-Al.
                                                                                                                                                                                                                                                                                                                                       ADC69646 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003194770-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC48535 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194773-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #205.
182003194776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD11102 gtandard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194774-A1.
99.5%;
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08-MAY-2003.
(GETH ) GENENTECH INC.
MATCh ... 13-rity 99.68;
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16-007-2003.
(GETH ) GENENTECH INC.
MAtch '-'arity 99.5%;
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(GETH ) GENENTECH INC.
Pry Match
99.5%;
st Local Similarity 99.6%;
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ry Match 99.5%;
t Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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Local Similarity 99.6%;
                                                                                                                         (GETH ) GENENTECH INC.
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              Best Local Similarity RESULT 134
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Best Local Similarity
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   Query Match
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Pred. No. 4.1e-114;
                  Length 229;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC54629 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087363-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC53590 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087364-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC59113 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC55991 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087360-A1.
                                                                  ADC57338 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087366-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADCS6561 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410
US2003087346-A1.
                                                                                                                                                                                                          ADC66529 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087367-A1.
                                                                                                                                                                                                                                                                                                                                         Novel human secreted and transmembrane protein, PRO733 US2001087141-41.
                  Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human PRO polypeptide #205.
US2003087362-A1.
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RESULT 128
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                99.5%;
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Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                     08-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 131
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Best Local Similarity
RESULT 132
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Best Local Similarity
RESULT 130
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              Query Match
Best Local Similarity
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                                                                                                                                                          Query Match
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RESULT 129

Length 229;

Length 229;

Length 229;

Length 229;

Length 229;

DB 7;

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Query Match
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     RESULT 152
1D ADD542
DE Novel
PN US2003
PD 30-OCT
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Query Match

99.5%; Score 1171; DB 7; Length 229;

Best Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 150
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 229;
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                                    ADC80043 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087358-A1.
                                                                                                                                                                                                                                                                                                                         ADD41225 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003203438-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aurosios standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003203437-A1.
   99.6%; Pred. No. 4.1e-114;
                                                                                                                              Score 1171; DB 7;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                       Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                                                                                              ADD09512 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194775-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD52364 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194769-A1.
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30-Orm Acceptance #205.
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Human PRO polypeptide #205.
US2003194779-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD02611 standard; protein; 229 AA. Human PRO polypeptide #205. US2003203431-A1.
                                                                                                                                                                                                                US20011...
16-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; SC
                                                                       US2005uc...
08-MAY-2003.
(GETH ) GENENTECH INC.
99.5%;
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A (GETH) GENENTECH INC.
OLARY MATCH 99.5%;
Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 148
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                                                                                                                          Query Match
Best Local Similarity
RESULT 144
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
Best Local Similarity
RESULT 143
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Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                  99.5%; Score 1171; DB 7; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                Length 229;
nuusquz7 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003203432-A1.
30-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB12351 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194765-A1.
                                                                                              Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                             LT 153
ADB49481 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003096744-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT.160
ID ADE35535 standard; protein; 229 AA.
DE Human secreted/transmembrane protein, PRO733.
PN US2003203434-A1.
                                                                                                                                                                                                                                                                                ADD92544 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199030-A1.
23-OCT-2003.
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Human PRO polypeptide #205.
US2003199057-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                   ADD91440 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE22283 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199056-A1.
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Best Local Similarity 99.6%;
                                                                                                                                                                                               22-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
Query Match
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                                                                                                                                                                                                                                          Best Local Similarity
RESULT 154
                                                                                                           Best Local Similarity RESULT 153
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Best Local Similarity
RESULT 157
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RESULT 161
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Length 1229;

Bast Local Similarity 99.5%; Score 1171; DB 7; Length 229;

Bast Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 164

ID ADE17860 standard; protein; 229 AA.

PN US2003199023-A1.

PD 23-OCT-20^2

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                                                                                         Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003194767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE34007 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194791-A1.
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4.1e-114;
                                                                                       Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
ADE16649 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203435-A1.
                                                                                                                                          ADD72264 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 1171;
Pred. No. 4.
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Human PRO polypeptide #205.
US2003199053-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD80059 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207417-A1.
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                                             PD 30-0CT-2003.
PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 99.6%;
                                                                                                                                                                                       PD 30-OCT-2003.

PA (GETH ) GENENTECH INC.

QUETY MARCH

Best Local Similarity 99.6%;

RESULT 163
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2003.
2003.
21Y Match
Best Local Similarity >> RESULT 165
ID ADD91992 stand-DE Human PROFON PN US20031
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RESULT 166
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Best Local Similarity
RESULT 170
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Best Local Similarity
RESULT 169
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(GETH ) GENENTECH INC.
Query Match 99.5%; Score 1171; DB 7; Length 229;
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003194766-A1.
                                                                        Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
Human secreted/transmembrane protein, PRO733. US2003194781-A1.
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Human secreted/transmembrane protein, PRO733.
                                                                                                                             ADE19516 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199025-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                           ADE43160 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199033-A1.
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Human PRO polypeptide #205.
US2003199059-A1.
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Human PRO polypeptide #205.
US2003199064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD78953 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003203429-A1.
                                                                                                                                                                                                                                                                             protein; 229 AA
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30-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; S'
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16-0CT-2003.
(GETH ) GENENTECH INC.
99.5$;
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16-007-2003.
(GETH ) GENENTECH INC.
*** MAtch ''...itv 99.6%;
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(GETH ) GENENTECH INC.
99.5%;
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(GETH) GENENTECH INC.
Query Match 99.5%;
Best Local Similarity 99.6%;
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Query Match
Best Local Similarity 99.6%;
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(GETH ) GENENTECH INC.
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st Local Similarity 99.6%;
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(GETH) GENENTECH INC.
10-GETY MATCh 99 5%;
Best Local Similarity 99.6%;
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Human PRO polypeptide #205.
US2003199032-A1.
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US2003199026-A1.
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06-NOV-2003
(GETH ) GEN
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Best Local S
                                                              RESULT 189
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                                          99.5%; Score 1171; DB 7; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003207384-A1.
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Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                   Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003195333-A1.
                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #205.
US2003199031-Al.
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Human PRO polypeptide #205.
US2003194777-A1.
                                                                                                                                                                                                                ADD89639 standard, protein, 229 AA. Human PRO polypeptide #205. US2003199028-A1.
                                                                                            protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE04722 standard; protein; 229 AA.
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Human NOVX4c protein.
WO2003050245-A2.
                                                                                                                                                                                                                                             US2001157...
23-OCT-2003.
(GETH ) GENENTECH INC.
Warch 99.5%; Sr
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23-OCT-2003,
(GETH ) GENENTECH INC,
99.5%; Sr
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(GETH ) GENENTECH INC.
99.5%; S.
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06-NOV-2003.
(GETH ) GENENTECH INC.
99.5%;
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RESULT 186
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                                                                                                         Human PRO polypeptide #205.
US2003207418-A1.
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US2003199034-A1.
           30-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                      Best Local Similarity
RESULT 180
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RESULT 181
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Best Local Similarity
RESULT 182
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RESULT 185
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                                                                                          ADD80611 standard;
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.US2003203433-A1.
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                                            Query Match
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Score 1171; DB 7; Length 229; Pred. No. 4.1e-114;
              Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207379-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH55340 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207381-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207385-A1.
06-NOV-2003.
            Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003206915-A1.
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Human secreted/transmembrane protein, PRO733.
US2003077700-A1.
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Human PRO polypeptide #205.
US2003207372-Al.
                                                             229 AA
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            99.5%;
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                                                     Human PRO polypeptide #205.
US2003207370-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 197
              Query Match
Best Local Similarity
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RESULT 190
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RESULT 192
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Best Local Similarity
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Length 229;

Length 229;

us-10-692-730-2.spdi.rag

RESULT 199

RESULT 201

PERCE

Query

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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                        ADC81147 standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US2003092115-A1.
99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003104536-A1.
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US20032110-A1.
                                                                                                                                                                                                                                                                          ADD76595 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003100087-A1.
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Human PRO polypeptide #205.
US2003092113-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE23387 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003092108-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE75811 standard; protein; 229 AA Human PRO polypeptide #205. US2003211571.A1. (GETH ) GENENTECH INC.
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15-MAY-2003.
(GETH ) GENENTECH INC.
99.5%;
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(GETH ) GENENTECH INC.

Query Match 99.5%;

Best Local Similarity 99.6%;
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Local Similarity 99.6%;
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Query Match 99.5%;
Best Local Similarity 99.6%;
RESULT 213
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(GETH ) GENENTECH INC.

Query Match 99.5%;

Best Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD86363 standard; protein;
Human PRO polypeptide #205.
US200320340-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 99.6%;
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Best Local Similarity 99.5%;
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Best Local Similarity 99.6%;
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US2003092111-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC.
Query Match
Best Local Similarity
RESULT 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE24582 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUN-2003
                                                                                                                                                                                                                                                                                                                                                           29-MAY-200
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                                                                                                                                                                                                                                                                                                                                                                                                                    Score 1171; DB 7; Length 229;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                       Length 229;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US22033087357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM82590 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733. US2003087355-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADNI5989 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087353-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADNI6618 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087385-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003087356-A1.
                                                                                                                                                                                                                                                                                          AD163559 standard; protein; 229 AA.
AD163559 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207387-A1.
06-NOV-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel human secreted and transmembrane protein PRO733. US2003207388-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH81421 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207377-A1.
                                                                                               ADI65060 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733 US2003207386-A1.
                                                                                                                                                                                                                            99.5%; Score 1171; DB 7; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                            Score 1171; DB 7;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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08-MAY-2003.
(GETH ) GENENTECH INC.
watch 99.5%; Sc
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08-MAY-2003.
(GETH ) GENENTECH INC.
40rch 99.5%; SC
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Best Local Similarity 9.

RESULT 206

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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
  (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 204
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RESULT 202
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Best Local Similarity
RESULT 203
                       Query Match
Best Local Similarity
RESULT 198
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Length 229;

DB 8; Length 229;

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Length 229;
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    Length 229;
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Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                        Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                             Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
    99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                           ADF46010 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195148-A1.
                                                ADF40214 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
23-00T-2003.
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102003203402-A1.
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                                                                                                                                                                                                                                                                                                       protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                             ADE91152 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003199061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE95293 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199052-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE93403 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199060-A1.
                                                                               US200315...
23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%; Si
                                                                                                                                                                                                         US200312-203.
16-007-2003.
(GETH ) GENENTECH INC.
99.5%;
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Local Similarity 99.6%;
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Best Local Similarity 99.6%;
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Local Similarity 99.6%;
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Best Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                ADE94741 standard; protein;
Human PRO polypeptide #205.
US2003199027-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
                Best Local Similarity
RESULT 222
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ID ADF24406 standard;
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      Query Match
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                                                                                                               Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                        Length 229;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                     99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADEB9882 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003130181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF61522 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195345-A1.
                                                                                                                                                       ADE89273 standard; protein; 229 AA. Human PRO polypeptide #205. US2003199062-A1. CGFTH ) GENENTECH INC.
                           ADD87407 standard; protein; 229 AA. Human PRO polypeptide #205. US2003203439-A1. 30-OCT-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                             ADE88721 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199054-A1.
                                                                                                                                                                                                                                                                                    ADE18412 standard; protein; 229 AA
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                                                                                                                            99.68;
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US2003194794-Al.
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(GETH ) GENENTECH INC.
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PANU J.
PAONI N F.
ROY M A.
SHELTON D L.
STEWART T A.
TUMAS D.
WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ASHKENAZI A J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GERRITSEN M E
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(GETH ) GENENTECH INC.
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KLJAVIN I J.
KUO S S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTSTEIN D. RESNOYERS L.
                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity
RESULT 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LVAROFF E
                                                                                                                         Best Local Similarity RESULT 217
                                                                                                                                                                                                                                                  Best Local Similarity RESULT 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
Best Local Similarity RESULT 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EATON D L.
FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GURNEY A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODOWSKI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GIRMALDI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-OCT-2003
                                                                                                             Query Match
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RESULT 220
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DB 8;

Query Match

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Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114;
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                                                                                                                      Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                           Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
   99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF27868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199437-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF33141 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003211091-A1.
                                                                                                                                                                                                                                                                                                         ADF27232 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF41462 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199435-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted/transmembrane protein, PRO733 US2003211092-A1.
                                  ADF33765 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003194780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    229 AA.
                                                                                                                                                                       ADF34984 standard; protein; 229 AA.
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                                                                                                                     99.5%;
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                                                                                                                                                                                       Human PRO polypeptide #205.
US2003199029-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                   16-OCT-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 231
ID ADF31765 standard; pp
DE Human secreted/transis
PN US2003194780-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC
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Best Local Similarity
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Best Local Similarity
RESULT 234
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                                                                                                                                   Best Local Similarity RESULT 232
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Best Local Similarity
RESULT 239
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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
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                                                                                                                                                                                                                                                             Length 229;
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                                                                                                                Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE91747 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003199058-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG22112 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207360-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG24305 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
                                                                                                                                                                                                                                     (GETH ) GENENTECH INC.
17 Match 99.5%; Score 1171; DB 8;
it Local Similarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
                                                                                                              Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
RESULT 240

ID ADF26608 standard; protein; 229 AA.

DE Human secreted/transmembrane protein, PRO733.

PN US2003199674-A1.

PD 23-OCT-2003.

PA (GETH) GENENTECH INC.
                                                                                                                                                                ADF34397 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003194410-A1.
                                                                                                                                                                                                                                                                                                                   ADF46634 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195344-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG20182 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207376-A1.
                                                     US2003150...
23-OCT-2003.
(GETH ) GENENTECH INC.
99.5%;
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RESULT 248
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Local Similarity 99.6%;
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ry Match 99.5%;
t Local Similarity 99.6%;
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Local Similarity 99.6%;
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Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG02326 standard; protein;
Human PRO polypeptide #205.
US2003207352-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 242
                                                                                                                        Best Local Similarity RESULT 241
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Best Local Similarity
RESULT 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 247
ID ADF98088 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2003207426-A1.
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Human PRO polypeptide #205. US2003207371-A1.

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Jest LOCATO 2017 252 3 ADG16796 standard; protect...

SULT 252 3 ADG16796 standard; protect...

B Human PRO polypeptide #205.

PN US2003207359-A1.

PD 06-NOV-2003.

PA (GETH ) GENENTECH INC.

99.5%; Score 1171; DB 8; Length 229;

""" Match "!" arity 99.6%; Pred. No. 4.1e-114;
                                                                            Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
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PA (CETH ) GENENTECH INC.
Query Match 99.5%; Score 1171; DB 8; Length 229;
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733
US2003207424-A1.
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(GETH) GENENTECH INC.
(GETH) 99.5%; Score 1171; DB 8;
ery Match
` `.`milarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
ADF98659 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003208055-A1.
                                                                                                                     ADG03490 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG19522 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207425-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG13359 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207357-A1.
                                                                                                                                                                                                                                                 protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG15586 standard; protein; 229 AA. Human PRO polypeptide #205. US2003219885-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99.54;
                                  PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 99.6%;
RESULT 250
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Human PRO polypeptide #205.
US2003207353-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 256
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Best Local Similarity
RESULT 258
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RESULT 254
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Best Local Similarity
RESULT 257
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06-NOV-2003.
(GETH ) GENENTECH INC.
ery Match 99.5%; Score 1171; DB 8; Length 229;
st Local Similarity 99.6%; Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003207427-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG07240 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207350-A1.
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US20032073356-Al.
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Novel human secreted and transmembrane protein PRO733.
US2003207390-A1.
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Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                 Novel human secreted and transmembrane protein PRO733 US2003207389-A1.
                                   Score 1171; DB 8;
Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Best Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 266
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(GETH) GENENTECH INC.
Query Match 99.5%; Score 1171; DB 8; Best Local Similarity 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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                                                                                                protein; 229 AA
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(GETH ) GENENTECH INC. 99.5%;
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Best Local Similarity 99.5%;
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1 (GETH ) GENENTECH INC.
10-CHT ) MATCH 99.5$;
BEST LOCAL Similarity 99.6$;
                                                                                            ADG06169 standard; protein;
Human PRO polypeptide #205.
US2003207374-A1.
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US2003207423-A1.
06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                                                            RESULT 259
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RESULT 260
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Length 229;

Score 1171; DB 8; Length 229; Pred. No. 4.1e-114; Novel human secreted and transmembrane protein PRO733. US2003207419-A1. ADG52455 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207414-A1. ADG54183 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733. US2003207416-A1. Score 1171; DB 8; Pred. No. 4.1e-114; Score 1171; DB 8; Pred. No. 4.1e-114; Score 1171; DB 8; Pred. No. 4.1e-114; Query Match
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 281 Score 1171; DB 8; Pred. No. 4.1e-114; אא 229 AA. Human secreted/transmembrane protein, PRO733. US2003215905-A1. Human secreted/transmembrane protein, PRO733. US2003216305-A1. ADG51868 standard; protein; 229 AA. Human secreted/transmembrane protein, PRO733. US2003215908-A1. ADG50620 standard; protein; 229 AA. Human secreted/transmembrane protein, PRO733 US2002207803-A1. (GETH ) GENENTECH INC. Human PRO polypeptide #205. US2003207805-A1. 06-NOV-2003. ADH30666 standard; protein; 229 AA. Human PRO polypeptide #205. US2003077723-A1. US2003.20-203. 20-NOV-2003. (GETH ) GENENTECH INC. 30-54; 99.54; S2003412. 20-NOV-2003. (GETH), GENENTECH INC. 99.5\$; US2003\_C. 06-NOV-2003. (GETH ) GENENTECH INC. Watch 99.5%; USZUUSZU 06-NOV-2003. (GETH ) GENENTECH INC. MATCh '17-11 99.5\$; PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 99.6%; 99.5%; 99.5%; 99.5%; (GETH ) GENENTECH INC. GENENTECH INC. 20-NOV-2003. (GETH ) GENENTECH INC. Best Local Similarity RESULT 279 Query Match Best Local Similarity Query Match Best Local Similarity Query Match Best Local Similarity Local Similarity Best Local Similarity 06-NOV-2003. (GETH ) GENE 06-NOV-2003 (GETH ) GEN 20-NOV-2003 24-APR-2003 Query Match Query Match Query Match RESULT 284 PD 06-NOV-2003.

PA (GETH) GENENTECH INC.

Query Match 99.5%; Score 1171; DB 8; Length 229;

Best Local Similarity 99.6%; Pred. No. 4.1e-114; 99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114; Score 1171; DB 8; Length 229; Pred. No. 4.1e-114; Score 1171; DB 8; Length 229; Pred. No. 4.1e-114; Length 229 Length 229 Length 229 Length 229 99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114; ADG58047 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207363-A1.
06-NOV-2003.
(GETH ) GENENTECH INC. ADG53631 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733. US2003207415-A1. ADG56943 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733. US2003207364-A1. ADGS8599 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733 US2003207368-A1. ADG70965 standard; protein; 229 AA. Novel human secreted and transmembrane protein PRO733. US2003207420-A1. ADG71517 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207421-A1. ADG57495 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207362-A1. ADG55839 standard, protein, 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207355-A1. Leary Match 10C.

Best Local Similarity 99.6%; Score 1171; DB 8; Len RESULT 275

ID ADG53631 standard; protein; 229 AA.

BE Novel human secreted and transparents of the post Score 1171; DB 8; Pred. No. 4.1e-114; Score 1171; DB 8; Pred. No. 4.1e-114; Score 1171; DB 8; Pred. No. 4.1e-114; Human PRO polypeptide #205. US2003207358-A1. US2003zv. ... 06-NOV-2003. (GETH ) GENENTECH INC. 99.5%; Sr US20032... 06-NOV-2003. (GETH ) GENENTECH INC. 99.5%; 99.5%; 99.5%; 99.5%; (GETH ) GENENTECH INC. (GETH ) GENENTECH INC. 06-NOV-2003. (GETH ) GENENTECH INC. 06-NOV-2003. (GETH ) GENENTECH INC. Best Local Similarity RESULT, 269 Local Similarity Best Local Similarity RESULT 272 Best Local Similarity RESULT 276 Best Local Similarity RESULT 268 Best\_Local\_Similarity RESULT 273 Best Local Similarity US2003207428-A1.

Length 229;

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Query Match

RESULT 271

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99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2003207366-A1.
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Novel human secreted and transmembrane protein PRO733
US2003207378-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207369-A1.
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Novel human secreted and transmembrane protein PRO733
US2003207429-A1.
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Pred. No. 4.1e-114;
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(GETH ) GENENTECH INC.
99.5%; Score 1171; DB 8;
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Pred. No. 4.1e-114;
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Human secreted/transmembrane protein, PRO733.
US2003216560-A1.
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US200302231-A1.
                                                 protein; 229 AA
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06-NOV-2003.
(GETH ) GENENTECH INC.
99.5%; Sr
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                                            ADG81152 standard; protein;
Human PRO polypeptide #205.
US2003194793-Al.
                                                                                                            (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 292
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Best Local Similarity
RESULT 289
Query Match
Best Local Similarity
RESULT 286
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RESULT 293
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Best Local Similarity
RESULT 294
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RESULT 287
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ADH25669 standard; protein; 229 AA.
Human neurotrimin homologue related protein sequence SBQ ID NO:447.
                                                                                                                             Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Novel human secreted and transmembrane protein PRO733.
US2004009548-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207382-A1.
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Novel human secreted and transmembrane protein PRO733
US2004009547-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207349-A1.
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Novel human secreted and transmembrane protein PRO733 US2003207383-A1.
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(GETH ) GENENTECH INC.

sry Match 99.5%; Score 1171; DB 8;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
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Pred. No. 4.1e-114;
              ADG59188 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
08-3AN-2004.
(GETH ) GENENTECH INC.
1267Y Match
171; DB
187Y Match
171; DB
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Human secreted/transmembrane protein, PRO733.
US2004006219-A1.
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Human PRO polypeptide #205.
US2003207361-A1.
06-NOV-2003.
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08-JAN-2004.
(GETH ) GENENTECH INC.
99.5%; S
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(GETH ) GENENTECH INC.
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Best Local Similarity RESULT 295
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 301
                                                                                                                                                                                                                                                                                                 Best
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Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                           99.5%; Score 1171; DB 8; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                    Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           / Match 99.5%; Score 1171; DB 8; Length 229; Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 1171; DB 8; Length 229;
Pred. No. 4.1e-114;
                                                                                                    Length 229;
nuvosoil standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2004039164 A1.
26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                    Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                  99.5%; Score 1171; DB 8; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JT 309
ADL07280 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004063921-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM17446 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004048332-A1.
                                                                                                                                                     ADJ77516 standard; protein; 229 AA. Human PRO polypeptide #205. US2004038336-Al. C6-FEB-2004 (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human PRO polypeptide #205.
US2003077659-A1.
                                                                                                                                                                                                                                                                                              ADJ65638 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004038335-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                 ADM27774 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004048333-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM42498 standard; protein; 229 AA. Human PRO polypeptide #205.
US2004058424-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99.5%;
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                                                                                                    99.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM28360 standard; protein;
Human PRO polypeptide #205.
US2004077064-A1.
                                                                                                                                                                                                                                                                                                                                                26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-MAR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-APR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 311
                                                                                                                                                                                                                                                        Best Local Similarity RESULT 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 309
                                                                                                                 Best Local Similarity RESULT 305
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S:
RESULT 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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(HINZ/) HINZMANN B.
(HINZ/) HINZMANN B.
(GAST) HERBANN K.
(CAST) HEIDEN CASTANOS-VELEZ E.
Query Match
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 1171; DB 8; Length 229; Pred. No. 4.1e-114;
                                                                                     Length 229;
                                                                                                                                                                                                                     Length 229;
                                                                                                                                                                                                                                                                                                                                                     Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                    ADU06673 standard; protein; 229 AA.
Novel bronchial cancer-associated human protein SegID899.
DE10316701-A1.
ID AD196394 standard; protein; 229 AA.

DE Novel human secreted and transmembrane protein PR0733.

PN US2003207354-A1.

PD 06-NOV-2003.

PA (GETH ) GENENTECH INC.

QUETY MATCH

PASEL Local Similarity 99.6%; Pred. No. 4.1e-114;
                                                                                                                                 ADS32346 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2004203125-A1.
                                                                                                                                                                                                                                                                                                             GETH) GENENTECH INC.

Query Match

Query Match

99.5%; Score 1171; DB 8;

Query Match

99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                     Score 1171; DB 8;
Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                               ADT03330 standard; protein; 229 AA. Human PRO polypeptide #205.
US2004214269-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADU50402 standard; protein; 229 AA PRO733, SEQ ID 447.
US2004223964-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 229 AA
                                                                                                                                                                       US20042...
14-0CT-2004.
(GETH ) GENENTECH INC.
*** Match '-'arity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 99.5%;
Local Similarity 99.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADW49681 standard, protein, PRO733 protein, SEQ ID 447.
US2005014226-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-NOV-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GERRITSEN M E. GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ASHKENAZI A J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GODOWSKI P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILVAROFF E. FONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SHELTON D L.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BOTSTEIN D.
DESNOYERS L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KLJAVIN I J
                                                                                                                                                                                                                             Best Local Similarity RESULT 315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KUO S S.
NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EATON D L.
FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PAN J.
PAONI N F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GRIMALDI J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HILLAN K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R M YOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAO W.
                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (TUMA/)
(WILL/)
(WOOD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DESN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FONG/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GODD/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GURN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HILL/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAON/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KUOS/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAPI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PANJ/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SHEL/
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FILVAROFF E.

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AEB14127 standard; protein; 229 AA.
Cancer cell diagnosis method-related human protein - SEQ ID 410.
US2005153396-A1.
                                                                                                                                                                                                                                          07-APR-2005.

1 (GETH) GENENTECH INC.

Query Match
99.5%; Score 1171; DB 9; Length 229;
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
 Length 229;
                                                                                                                                         Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99.5%; Score 1171; DB 9; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                           99.5%; Score 1171; DB 9; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99.5%; Score 1171; DB 9; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                              ADZ52342 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2005084935-A1.
                                                                                                                                                                                     ADZ03381 standard; protein; 229 AA.
Human secreted/transmembrane PRO733 protein.
US2005074837-A1.
                                            AEA99614 standard; protein; 229 AA.
Human PRO733 protein SEQ ID NO:2.
US2005118635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JUN-2005.

(GODD/) GODDARD A.

(GORN/) GODWSKI P J.

(SWIY/) GURNEY A L.

(SMIT/) SMITH V.

(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                 ASHKENAZI A J.
BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 5
RESULT 323
ID ABB14127 standard; prc
DE Cancer cell diagnosis
PD 14-JUL-2005.
PA (BREK) BAKER K P.
PA (BREK) BENESIMI M.
PA (BREK) BENESIMI M.
PA (BREK) BENESIMI M.
PA (BREK) BENESIMI M.
PA (BREN) DEFONCE I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GODOWSKI P J.
GRIMALDI J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PAN J.
PAONI N F.
ROY M A.
SHELTON D L.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TUMA/) TUMAS D.
(WILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
Query Match
Best Local Similarity
RESULT 319
                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                  DESNOYERS L.
EATON D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FERRARA N.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 322
                                                                                                                                                                                                                                                                                                                                                                                                                                  SOTSTEIN D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KLJAVIN I J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GURNEY A L.
HILLAN K J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               KUO S S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GAO W
                                                                                                                                                                       RESULT 320
                                                                                                                                                                                                                                                                                                            RESULT 321
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Human cancer-related PRO protein amino acid sequence - SEQ ID 410.
US2006040351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 99.5%; Score 1171; DB 10; Length 229;
Best Local Similarity 99.6%; Pred. No. 4.1e-114;
RESULT 327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-4PR-2006.
(GETH ) GENENTECH INC.
(ery Match 99.5%; Score 1171; DB 10; Length 229;
ery Match 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 1171; DB 10; Length 229; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                PD 13-OCT-2005.

PA (GETH ) GENENTECH INC.

Query Match

Query Match

Best Local Similarity 99.6%; Pred. No. 4.1e-114;

RESULT 325

ID AED86325 standard; protein; 229 AA.

DE Human PRO amino acid sequence, seq id 410.

PN US2005245730-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       99.5%; Score 1171; DB 9; Length 229; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                 Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AEH18401 standard; protein; 229 AA.
Human PRO protein amino acid sequence - SEQ ID 447.
US2006078964-A1.
                                                                                                                                                                                   99.5%; Score 1171; DB 9; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 99.6%; Fred. No. 4
RESULT 326
ID ARGS8272 standard; protein; 229 AA.
DE Human PRO733 polypeptide SEQ ID No: 410.
PD 06-APR-2006.
PA (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                 AED48170 standard; protein; 229 AA.
Human PRO733 amino acid sequence.
US2005227342-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AEI43917 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEI17335 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99.5%;
                         GURNEY A L.
GURNEY A L.
SHEWAND S.
SHITH V.
STEWART T A.
TUMAS D.
WATHWABE C K.
WOOD W I.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                    03-NOV-2005.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAO W.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SMITH V.
STEWART T A.
TUMAS D.
WATANABE C K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BAKER K P.
BERESINI M.
DEFORGE L.
DESNOYERS L.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GODDARD A. GODOWSKI P J.
                                                                                                                                                                                                Best Local Similarity RESULT 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GURNEY A L. SHERWOOD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZHANG Z.
                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                               (WATA/)
(WOOD/)
(ZHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BERE/)
                             (GODO/)
(GURN/)
(SHER/)
                                                                                                                   (TOMA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (DESN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SMIT/)
(STEW/)
                                                                                                  (STEW/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WATA/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GURN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GODD/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 329
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AEI17411 standard; protein; 229 AA.
Human cancer treatment biomarker amino acid sequence - SEQ ID 298.
US2006121511-A1.
Human cancer treatment biomarker amino acid sequence - SEQ ID 222.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 229;
                                                                                                                   Length 229;
                                                                                                                                                                                                                                                                                                                         Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USCOTATION OF THE SCI INC. (HUMA-) HUMAN GENOME SCI INC. (HUMA-) HUMAN GENOME SCI INC. 88.3%; Score 1039; DB 8; Length 456; ery Match 83.9%; Pred. No. 8.2e-100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99.0%; Score 1165; DB 7; Length 229; 99.1%; Pred. No. 1.8e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            wuery Match 70.9%; Score 835; DB 7; Length 193;
Best Local Similarity 93.1%; Pred. No. 6.4e-79;
RESULT 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE95518 standard; protein; 162 AA.
Human NOVX4b protein.
WQ2003050245-A2.
19-JUN-2003.
(CURA-) CURAGEN CORP.
70.3%; Score 828; DB 7; Length 162;
                                                                                                                                                                                                                                                                                                                                                                                    AEI24000 standard; protein; 229 AA.
Human secreted/transmembrane protein PRO733, SEQ ID NO:410.
EP1672070-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY35962 standard; protein; 229 AA.
Extended human secreted protein sequence, SEQ ID NO. 211
WO9931236-A2.
                                                                                                                                                                                                                                                                                                                         99.5%; Score 1171; DB 10; 99.6%; Pred. No. 4.1e-114;
                                                                                                                     99.5%; Score 1171; DB 10; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99.5%; Score 1171; DB 10; 99.6%; Pred. No. 4.1e-114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99.1%; Score 1166; DB 2; 99.1%; Pred. No. 1.4e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99.1%; Score 1166; DB 8; 99.1%; Pred. No. 1.4e-113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADR03639 standard; protein; 456 AA.
Human T1/ST2 receptor binding protein.
US2004146501-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP19270 standard; protein; 229 AA. Human secreted polypeptide #121. US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 334
ID ADE95516 standard; protein; 229 AA.
DE Human NOVX4a protein.
PN W02003056245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG14988 standard; protein; 193 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human SECP-21 protein.
WO2003087300-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
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(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-OCT-2003.
(INCY-) INCYTE CORP.
                                   08-JUN-2006.
(LEEH/) LEE H.
(SHAW/) SHAW P.
(CLAR/) CLARK E A.
                                                                                                                                                                                                                                          PD 08-JUN-2006.
PA (LEEH/) LEE H.
PA (SHAW/) SHAW P.
PA (CLARK E A.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match
Best Local Similarity
RESULT 333
                                                                                                                                   Best Local Similarity RESULT 330
                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 331
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GEST ) GENSET.
                    US2006121511-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-JUN-2006
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AULZ2684 standard; protein; 141 AA.
Human disease detection and treatment (MDDT) protein - SEQ ID 133.
WO2003062379-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO23233 standard; protein; 169 AA.
Human breast tumour associated protein 47-like polypeptide NOV1
US2003027158-A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vuery Match
Best Local Similarity 97.5%; Pred. No. 9.6e-76;
RESULT 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 804; DB 7; Length 169;
Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 31-JUL-2003.

A (INCY-) INCYTE GENOMICS INC.

Query Match 57.1%; Score 672; DB 7; Length 141;

Best Local Similarity 99.3%; Pred. No. 5.4e-62;
                                                                                                                                                                                                                                                                                                       69.8%; Score 821; DB 8; Length 162; 98.1%; Pred. No. 1.5e-77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             68.3%; Score 804; DB 4; Length 169; 97.5%; Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 169;
                                                                                                                                           69.8%; Score 821; DB 2; Length 162; 98.1%; Pred. No. 1.5e-77;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 169;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 169;
                                         AAY35900 standard; protein; 162 AA.
Extended human secreted protein sequence, SEQ ID NO. 149.
W09931236-A2.
24-JUN-1999.
(GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 804; DB 8;
Pred. No. 9.6e-76;
Best Local Similarity 98.8%; Pred. No. 2.7e-78; RESULT 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           68.3%; Score 804; DB 6; 97.5%; Pred. No. 9.6e-76;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 804; DB 8;
Pred. No. 9.6e-76;
                                                                                                                                                                                RESULT 339
ID ADP19208 standard; protein; 162 AA.
DE Human secreted polypeptide #59.
PN US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                            AAB61129 standard; protein; 169 AA.
Human NOV1 protein.
WQ200075321-A2.
14-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADMS6365 standard; protein; 169 AA.
Human cell adhesion molecule NOV1.
US2003082554-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADF66729 standard; protein; 169 AA.
Novel human protein NOV1.
US2003199103-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADI19766 standard; protein; 169 AA. Human NOV1 protein.
US2004002134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AD060239 standard; protein; 169 AA.
Human NOV1 protein.
US2003134430-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              68.3%;
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97.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 68.3%;
Local Similarity 97.5%;
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(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-JUL-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-OCT-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-2003.
(CURA-) CURAGEN CORP.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 346
ID ADL22684 standard;
                                                                                                                                                                                                                                                              10-JUN-2004.
(GEST ) GENSET SA.
                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Length 227;

Length 133;

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RESULT 356

ID AEB81004 standard, protein; 227 AA.

DE Human putative TI/STZ receptor binding protein precursor SEQ ID NO:3.

PD 19205287606-A1.

PD 29-DEC-2005.

PA (HUMA) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEMY ) GENETICS INST INC.
52.3%; Score 615; DB 3; Length 133;
it Local Similarity 96.8%; Pred. No. 4.9e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43.0%; Score 506; DB 2; Length 120; 100.0%; Pred. No. 1.2e-44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 317; DB 2; Length 61;
Pred. No. 3.2e-25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY35901 standard; protein; 120 AA.
Extended human secreted protein sequence, SEQ ID NO. 150.
WO9931236-A2.
                                                                                                                                                                                                      AAWS4441 standard; protein; 133 AA.
Mouse novel secreted protein isolated from clone AI6_li
WO9814470-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABB64192 standard; protein; 235 AA.
Drosophila melanogaster polypeptide SEQ ID NO 19368
WO200171042-A2.
                                                                                                                                               DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 416.5; DB 4;
Pred. No. 7.9e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            43.0%; Score 506; DB 8; I 100.0%; Pred. No. 1.2e-44;
                                                                                                                                                                                                                                                                                                            Score 615; DB 2;
Pred. No. 4.9e-56;
                                                                                                                                            52.3%; Score 615.5; DB 1
55.0%; Pred. No. 9.3e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52.1%; Score 613.5; DB 5 54.5%; Pred. No. 1.6e-55;
      Best Local Similarity 56.7%; Pred. No. 8.1e-56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         39.8%; Score 469; DB 5; 91.1%; Pred. No. 7.8e-41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABP41689 standard; protein; 240 AA.
Human ovarian antigen HE2DQ62, SEQ ID NO:2821.
WO200200677-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 364
ID AAY12211 standard; protein; 61 AA.
DE Human 5' EST secreted protein SEQ ID NO: 524.
PN W09906554-A2.
                                                                                                                                                                                                                                                                                                                                                                AAB10248 standard; protein; 133 AA.
Human adult blood protein fragment AI6_1i.
WO200037630-A1.
29-JUN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADP19209 standard; protein; 120 AA. Human secreted polypeptide #60. US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK35641 standard; protein; 107 AA.
Novel human polypeptide SegID7723.
WO200216439-A2.
                                                                                                   29-DEC-2005.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                        WOSLE--
(GEMY ) GENETICS INST INC.
(GEMY MATCh 52.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26.9%;
98.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   35.4%;
                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                             Best Local Similarity RESULT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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(GEST ) GENSET SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JUN-1999
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                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEE81034 standard; protein; 227 AA.
Human putative T1/ST2 receptor binding protein precursor SEQ ID NO:3.
US2005287606-A1.
                AAU78099 standard; protein; 227 AA.
Human T1/ST2 receptor binding protein (ILIRLILG) variant sequence.
WO200216646-A1.
28-FBB-2002.
                                                                                                                                                                                                                                                                                                                                           AAU78098 standard; protein; 227 AA.
Human TI/ST2 receptor binding protein (ILIRLILG) protein sequence.
WO200216646-A1.
28-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 10; Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 7; Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               53.2%; Score 626.5; DB 7; Length 227; 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                     Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53.2%; Score 626.5; DB 8; Length 227; 55.9%; Pred. No. 6.5e-57;
                                                                                                                        Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 125;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 2; Length 224;
                                                                                                                                                                                                                                                                                 53.2%; Score 626.5; DB 2; 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                          (GENA-) GENAISSANCE PHARM INC.
ry Match
t Local Similarity 55.9%; Pred. No. 6.5e-57;
                                                                                           (GENA.) GENAISSANCE PHARM INC.
ry Match
t Local Similarity 56.3%; Pred. No. 9.4e-58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-02-2
29-02-2
(HUMA-) HUMAN GENOME SCI INC.
(ery Match
62-5; DB 10
67; Arilarity 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JUL-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
ery Match 52.9%; Score 623; DB 8;
ery Match 99.2%; Pred. No. 6.4e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          53.2%; Score 626.5; DB 7 55.9%; Pred. No. 6.5e-57;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB87501 standard; protein; 227 AA.
Human T1 receptor (T1R) ligand polypeptide.
US6586210-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) 19-NOV-1996.
\ (IMMV) IMMUNEX CORP.
$2.3%; Score 616;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW09640 standard; protein; 224 AA.
Murine cytokine, ST2 ligand polypeptide
US5576191-A.
                                                                                                                                                                            AAW00639 standard; protein; 227 AA.
Human cytokine, ST2 ligand polypeptide.
US5576191-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADN35149 standard; protein; 227 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADQ66619 standard; protein; 125 AA.
Novel human protein sequence #1592.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI34090 standard; protein; 227 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-OCT-2003. (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human Tl receptor ligand.
US2003100048-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human TlR ligand protein.
US2003198618-Al.
                                               Loos.

Lory Match
Best Local Similarity
RESULT 348
ID AAW09639 stand>
DE Human cyto*
PN US55761
                                                                                                                                                                                                                                          19-NOV-1996.
(IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 353
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Best Local Similarity
RESULT 355
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NIJJ/) NI J.
(GENT/) GENTZ R.
(RUBE/) RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JUL-2003
                                                                                                                                                                                                                                                                                   Query Match
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RESULT 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 351
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Length 240;

Length 107;

Length

Length 120;

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ABU96190 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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                                                                                        Query Match
Best Local Similarity
RESULT 375
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                                                                                                                                                                                                                                                                              ABU84324 standard;
                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                         06-FEB-2003.
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RESULT 374
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 10; Length 224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23.9%; Score 281.5; DB 10; Length 224; 32.7%; Pred. No. 1.1e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INCYTE GENOMICS INC.

23.9%; Score 281.5; DB 6; Length 231;
t Local Similarity 32.7%; Pred. No. 1.2e-20;
                                                                                                                                                                                                                                                            DB 4; Length 224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 22.6%; Score 266.5; DB 2; Length 217; Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                          Length 215;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 219;
                                                                                                                                                                                                                                                                                                                                                                                                             23.9%; Score 281.5; DB 8; Length 224; 32.7%; Pred. No. 1.1e-20;
                                                                                                                                                                                                                                                                                                   AULJ1890 standard; protein; 224 AA.
Human protein encoded by a full length cDNA clone SeqID 3923.
EP1396543-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW37863 standard; protein; 217 AA.
Amino acid sequence of the human type-1 membrane protein
WO9811215-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABB11780 standard; peptide; 219 AA.
Human T1 receptor ligand III homologue, SEQ ID NO:2150
WO200157188-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23.7%; Score 278.5; DB 4; 32.3%; Pred. No. 2.3e-20;
                                                               G3-SEP-1998.

(HUWA-) HUMAN GENOME SCI INC.
24.2%; Score 284.5; DB 2;
ery Match
-1 cimilarity 32.4%; Pred. No. 5.2e-21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEK00714 standard; protein; 224 AA.
Protein aggregation-related protein SEQ ID NO:8.
WO2006091964-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23.9%; Score 281.5; DB 1 32.7%; Pred. No. 1.1e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 280.5; DB 4
Pred. No. 1.4e-20;
                                                                                                                                                                                                                                                            23.9%; Score 281.5; DB 4 32.7%; Pred. No. 1.1e-20;
                 AAW70459 standard; protein; 215 AA.
Human T1-receptor ligand III splice variant 2.
WO9838311-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB92820 standard, protein; 224 AA.
Human protein sequence SEQ ID NO:11348
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEL84842 standard; protein; 224 AA.
Tumor marker gene TMED7 SEQ ID NO 209.
WO2006110593-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR41674 standard; protein; 231 AA.
Human DITHP cell membrane protein.
WO200297031-A2.
                                                                                                                                                              AAW93844 standard, protein; 224 AA.
Human polypeptide, SEQ ID NO: 3923.
EPI130094-42.
(HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                       10-MAR-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SAGA ) SAGAMI CHEM RES CENTRE. (PROT-) PROTEGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 32.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                              A.S. 2004.
A.S.) RES ASSOC B.
A.S.Y MATCh
Best Local Similarity
RESULT 368
ID AEK00714 stand*
DB Protein acr
PN W020066*
PD 31
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2006.

Au-) UNIV ALABAN.

LY Match
Best Local Similarity ...
RESULT 369
ID AEL84842 standar
DD Tumor markar
PN W020061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYAL-) UNIV ALABAMA.
Query Match
                                                                                                            Query Match
Best Local Similarity
RESULT 366
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Best Local Similarity
RESULT 367
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-DEC-2007
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RESULT 365
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RESULT

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                      6; Length 217;
                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                            22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 217;
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                                                                                                                                                                                               Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABJ37053 standard; protein; 217 AA.
Human breast cancer / ovarian cancer related protein #29
WO2003000012-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU89888 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2001036147-A1.
20-FEB-2003.
                                                                                                                       ABU88009 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003032127-A1.
13-FBE-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted polypeptide PRO1078, SEQ ID NO:124. US:2003036159-Al.
                                                                                                                                                                                                                                                                                                                                                 Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003027278-A1.
                                                                                                                                                                                               Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR68137 standard, protein, 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027264-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                    Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein (PRO) #62.
US2003040070-A1.
27-FEB-2003.
                                                                                                                                                                                                                                                                  Human secreted/transmembrane protein (PRO) #62.
US2003032112-A1.
13-FEB-2003.
ABU58461 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003027272-A1.
                                                                                                                                                                                                                                                   protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABU82767 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003032113-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JAN-2003.
(MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                               22.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22.6%;
31.6%;
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31.6%;
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31.6%;
                                                                      22.6%;
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31.6%;
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Length 217;

Length 217;

DB 6;

23

us-10-692-730-2.spdi.rag

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                  Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU85024 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
                                                                                                                                                                Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR98803 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR92226 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036160-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR78288 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054474-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR99413 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040063-A1.
                                                                                                                                                                                                                                                                                                        Score 266.5; DB 6 Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                               ABU67467 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036162-A1.
                                                                                         Human secreted/transmembrane protein (PRO) #62. US2003036146-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO16326 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003027267-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO18867 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044925-A1.
                                                                                                                                                                                                                                                                                                                                                         ABU80495 standard; protein; 217 AA. Human PRO protein #62. US2003036137-A1.
                                                                          ABU86254 standard; protein; 217 AA
                                                                                                         US2003002.
20-FEB-2003.
(GETH ) GENENTECH INC.
22.6%; S
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20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S.
                                                                                                                                                                                                                                                                                                                                                                                          US20030502.
20-FEB-2003.
(GETH ) GENENTECH INC.
22.6%;
                    22.6%;
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31.6%;
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                                                                                                                                                                                                                                                                                                                         Local Similarity
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Best Local Similarity
                  Query Match
Best Local Similarity
RESULT 395
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Best Local Similarity
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     20-FEB-2003.
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                                                                                                                                                                                                                                                                       20-FEB-2003
                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                           RESULT 397
                                                                                                                                                                                                  RESULT 396
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                                                                                                                                                                                                                                                                                                                          Best
                                                                                                                                                                            Length 217;
                                                      Length 217;
                                                                                                                                                                                                                                                                                                      Length 217;
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Pred. No. 4.1e-19;
Novel human secreted and transmembrane protein PRO1078.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABU98799 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003013153-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABU98014 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003017544-A1.
23-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABU91720 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003027277-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR94666 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003044926-A1.
                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                              Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                      DB 6;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040056-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                      Human secreted/transmembrane protein (PRO) #62.
US2003036149-A1.
20-FEB-2003.
                                                                                                                                                                                                                                              Human secreted/transmembrane protein (PRO) #62.
US2003044923-A1.
06-MAR-2003.
                                                                                                                                                                                                                                                                                                                                             Human secreted/transmembrane protein (PRO) #62.
US2003040062-A1.
27-FEB-2003.
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU85639 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003036140-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #62.
US2003036141-A1.
                                                                                                         protein; 217 AA
                                                                                                                                                                                                                                 protein; 217 AA
                                                    22.6%;
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 391
                                         Query Match
Best Local Similarity
RESULT 385
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Best Local Similarity
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Best Local Similarity
RESULT 394
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Best Local Similarity
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Best Local Similarity
RESULT 393
                                                                                                       ABU92621 standard;
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                US2003036144-A1.
                                20-FEB-2003.
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RESULT 388

BEE

2 2 E 1

RESULT 389

2282

RESULT 390

A B E B B

RESULT 392

2 2 E C

DB 6; Length 217;

Length 217;

9

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Query Match
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                                  22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                     Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                    Length 217;
                                                       Novel human secreted and transmembrane protein PRO1078.
US2003032101-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO06210 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003022294-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           ABU8B714 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003036133-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR59246 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027275-A1.
                                                                                                                                                    DB 6;
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(GETH ) GENENTECH INC.
(ery Match 22.6%; Score 266.5; DB 6;
ery Match 31.6%; Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003036123-A1.
20-FEB-2003.
                                                                                                                                                    22.6%; Score 266.5; DB 6 31.6%; Pred. No. 4.1e-19;
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US2003027324-Al.
                                                                                                                                                                                                                                                                                                                        ABO02140 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003040054-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU83409 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036134-A1.
                                                                                                                                                                                                                   Human secreted/transmembrane protein (PRO) #62
US2003036124-A1.
                                                                                                                                                                                                     ABO11495 standard; protein; 217 AA
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Luda Match
Best Local Similarity 3.
RESULT 414
I DE Human ser
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cEB-2003.
cary Match
Best Local Similarity or RESULT 413
ID ABO19172 stand or DE Novel humar
PN US2003.
PD 20.
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US2003032114-A1.
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Best Local
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                       22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR67113 standard, protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027266-A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                       ÄBR66808 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US20030148-A1.
20-PEB-2003.
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein, SEQ ID 124
US200303036156-A1.
20-FEB-2003.
                                                                                                                                                                                  Human secreted/transmembrane protein (PRO) #62. US2002040060-A1.
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Human secreted/transmembrane protein (PRO) #62.
US2003036128-A1.
                                                                                                                                                                                                                                                                                                                                       ABO13727 standard, protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US203004916-A1.
06-MAR-2003.
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Human secreted/transmembrane protein (PRO) #62.
US2003054483-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABUSS997 standard; protein; 217 AA.
Human secreted/transmembrane protein, PRO1078.
US2003022298-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO07478 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003032117-A1.
13-FEB-2003.
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31.6%;
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31.6%;
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A (GETH) GENENTECH INC.
A LOURY MATCH 22.6%;
Best Local Similarity 31.6%;
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Query Match
Best Local Similarity
RESULT 415
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Best Local Similarity
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Best
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                                                                           Length 217;
                                                                                                                                                                                Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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          ABU95270 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003036117-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                   ABR70024 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003032138-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR69357 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003035132-A1.
20-FEB-2003.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
18200332137-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR67832 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003027269-A1.
                                                                         22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                         DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR65220 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027268-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                DB 6;
                                                                                                                                                                                                                                                                                         Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                  ABU71173 standard; protein; 217 AA. Human PRO1078 protein. US200305143-A1. 20-FEB-2003.
                                                                                                                                                                                                                              protein, 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU81300 standard; protein; 217 AA. Human PRO polypeptide #62. US2003017542-Al. 23-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AB001498 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003008353-A1.
                                                                                                                                                                                22.6%;
                                                                                                                                                                                                                                                                                        22.6%;
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
RESULT 429
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31.6%;
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PA (GETH ) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;

RESULT 431
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31.6%;
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31.6%;
                                                                                                                                                                                                                         AB007783 standard, protein
Human PRO polypeptide #62.
US2003032130-Al.
13-FEB-2003.
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u03036117-A1.

20-FEB-2003.

Query Match

Best Local Similarity -

RESULT 426

ID ABU71173 stand

DE Human PRCT

PN US200
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Best Local Similarity
RESULT 430
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Best Local Similarity
RESULT 428
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Best Local Similarity
RESULT 434
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Best Local Similarity
RESULT 432
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Best Local Similarity
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RESULT 435
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                                                                                                                                                                                Query Match
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RESULT 425
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DB 6; Length 217;
                                                                                                                                                                                                                                              DB 6; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABU94960 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003032123-A1.
13-PEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU90508 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003032108-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABU93670 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003032119-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR64915 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003027263-A1.
ABR68442 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027274-A1.
06-FRB-2003.
                                                                                                                                      ABR71854 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032135-A1.
                                                                                                                                                                                                                                              Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                  Score 266.5; DB 6
Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003022297-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU83104 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032105-A1.
13-FEB-2003.
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Human secreted/transmembrane protein (PRO) #62
US2003032111-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                        protein; 217 AA.
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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Best Local Similarity 31.6%;
RESULT 437
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Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                              Query Match
Best Local Similarity
RESULT 436
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Best Local Similarity
RESULT 438
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Best Local Similarity
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Best Local Similarity
RESULT 456
                                                                                                                       Best Local Similarity RESULT 457
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Best Local Similarity
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30-JAN-2003.
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       Query Match
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                                                  Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                          Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                         DB 6; Length 217;
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Novel human secreted and transmembrane protein PRO1078.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                  ABU85944 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US20030223300-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU82231 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003036136-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU81799 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
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Pred. No. 4.1e-19;
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9
Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003027271-A1. 06-FEB-2003.
                                                                                                                                                                                           Human secreted polypeptide PRO1078, SEQ ID NO:124
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Pred. No. 4.1e-19;
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Pred, No. 4.1e-19;
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31.6%; Pred. No. 4.1e-19;
                                                                                                          Human secreted/transmembrane protein (PRO) #62.
US2003036125-A1.
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US2003032109-A1.
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27-PFB-2003
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US2003027280-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABU87242 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003036138-A1.
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                                                                                              protein; 217 AA
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31.6%;
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RESULT 455
                                                             Best Local Similarity RESULT 446
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RESULT 454
                                                                                              ABO06563 standard;
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
6; Length 217;
                                                                                                                                          22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                    22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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Novel human secreted and transmembrane protein PRO1078.
US2003032131-A1.
                                                                                                                                                                                                   ABU93980 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003036155-A1.
20-FBB-2003.
                                                                                                                                                                                                                                                                                                                                             ABU99833 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-FEB-ZUUJ.
(GETH ) GENENTECH INC.
227 Match 22.6%; Score 266.5; DB 6;
Trans Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR66503 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027281-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                        ABRS9792 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032120-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR90921 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
US2003040058-A1.
27-FEB-2003.
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
  22.6%; Score 266.5; DB 6, 31.6%; Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003032129-A1.
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US2003017540-Al.
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31.6%;
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31.6%;
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31.6%;
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31.6%;
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US2003032106-A1.
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US2003032103-A1.
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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                     ABR65893 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003036165-A1.
                                                                                                        ABR70329 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032139-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                               ABR64610 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027262-A1.
                                                                                                                                                                                                                                                                DB 6;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003036142-A1.
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     Human PRO polypeptide #62.
US2003032107-Al.
13-FEB-2003.
                                                                                                                                                                                                          protein; 217 AA.
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US2003036145-Al.
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20-FEB-2003.
(GETH ) GENENTECH INC.
Warch 22.6%; St
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31.6%;
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31.6%;
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US2003022301-A1.
30-JAN-2003.
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Best Local Similarity
RESULT 470
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RESULT 474
                                                                             Best Local Similarity RESULT 467
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RESULT 469
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Best Local Similarity
RESULT 471
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Best Local Similarity
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                                                                                                                                                                                                         ABU98494 standard;
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RESULT 466
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Length 217;
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                                                                                                                                                                Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US200303212-A1.
13-FEB-2003.
                                                DB 6;
                                                                                                                                                                                                              ABR70919 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040069-A1.
27-FEB-2003.
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Pred. No. 4.1e-19;
                                               Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
Human secreted/transmembrane protein (PRO) #62.
US2003044931-A1.
06-MAR-2003.
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Pred. No. 4.1e-19;
                                                                                       Human secreted/transmembrane protein (PRO) #62.
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Human secreted/transmembrane protein (PRO) #62
US2003032116-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO09003 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003036152-A1.
20-FEB-2003.
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Human secreted/transmembrane protein (PRO) #62
US2003017543-A1.
                                                                                                                                                                                                                                                                                                                             ABU87547 standard; protein; 217 AA. Human PRO polypeptide #62.
US2003022293-A1.
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US2003036139-A1.
20-PEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                protein; 217 AA
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31.6%;
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Human PRO polypeptide #62.
US2003032128-A1.
13-PEB-2003.
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Human PRO polypeptide #62.
US2003017541-A1.
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                                                              Best Local Similarity RESULT 477
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Best Local Similarity
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Best Local Similarity
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                                                Query Match
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13-MAR-2003
(GETH ) GEN
 22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                               22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                         Length 217;
                                                                                                                                                                                                                                                                                                                               ABU96789 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003032140-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO04985 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003008352-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049741-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR70634 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003040076-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003036135-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054455-A1.
20-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                         22.6%; Score 266.5; DB·6; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003044922-A1.
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US2003032118-A1.
                                                      ABUI0571 standard; protein; 217 AA.
Human secreted/transmembrane protein #62.
US2002127584-A1.
                                                                                                                                                                                                  ABU95580 standard; protein; 217 AA. Human PRO polypeptide #62. US2003032115-A1.
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27-FEB-2003.
(GETH ) GENENTECH INC.
"orch 22.6%; SC
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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RESULT 490
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RESULT 491
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RESULT 492
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Query Match
Best Local Similarity
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                                       RESULT 487
                                                                                                                                                                                       RESULT 488
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PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PD 10-APR-2003.
PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 217;
                 Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                 Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                     Length 217;
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                                                                                                                                                                                                        ADMINUTY Standard; protein; 217 AA. Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                           ABR88481 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068743-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM77302 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054479-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM07948 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068752-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                    Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6;
                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO31531 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068725-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AB028786 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003068685-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO40428 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068682-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO35853 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068701-A1.
10-APR-2003.
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13-WRR-2003.
(GETH ) GENENTECH INC.
Match '''arity 31.6%;
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%;
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20-MMR-2003.
(GETH ) GENENTECH INC.
Match ''rrity 31.6%;
                                                                                                                                                               Best Local Similarity 31.6%;
RESULT 498
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(GETH ) GENENTECH INC.
22.6%;
                 Query Match 22.6%;
Best Local Similarity 31.6%;
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                                                                                                                              13-MAR-2003.
(GETH ) GENENTECH INC.
GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
RESULT 500
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Best Local Similarity
RESULT 501
                                                    RESULT 497
ID ABR81183 standard;
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RESULT 499
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                                            Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                            Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
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                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM17225 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054459-A1.
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20.MAR-2003.

(GETH ) GENENTECH INC.

(GETY Match 22.6%; Score 266.5; DB 6;

(ery Match 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR95276 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040071-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 6;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR94971 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044930-A1.
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted/transmembrane protein (PRO) #62.
US2003036131-A1.
                                                                                                                                                                                                                    Human secreted/transmembrane protein (PRO) #62 US2003073180-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO21514 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054471-Al.
                                                                                                                                                                                                     ADA77876 standard; protein; 217 AA
                                                                                    protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR97778 standard; protein; 217 AA
                                                                                                                                                                                                                              US20030,...
17-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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27-FEB-2003.
(GETH ) GENENFECH INC.
22.6%; S/
                                                                                                Human PRO polypeptide #62.
US2003068755-A1.
10-APR-2003.
(GETH ) GENENTECH INC.
Duery Match 22.6%; Steet Local Similarity 31.6%; P.
(GETH ) GENENTECH INC. 22.6%;
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31.6%;
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                                                     Best Local Similarity
RESULT 506
                                                                                                                                                                      Best Local Similarity RESULT 507
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Best Local Similarity
RESULT 513
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RESULT 508
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RESULT 509
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Best Local Similarity
RESULT 511
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RESULT 514
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                                                                                    ABO43992 standard;
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                                            Query Match
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Length 217;
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                                                              Length 217;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064452-A1.
                                                                                                          ABR87566 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068705-A1.
                                                                                                                                                                                                                              ABM77607 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054473-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM06118 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068704-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM03624 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068722-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM35075 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073183-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR92836 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
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                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                            9
                                                                                                                                                                                                                                                                                                                                                     ABM27837 standard, protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
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                                                                                                                                                                                     Score 266.5; DB 6 Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                              Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
               US200300...
03-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
                                                                                                                                                                                                                                                          US200305...
20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
                                                                                                                                              PD 10-ARR-2003.
PA (GETH ) GENENTECH INC.
Query March 22.6%;
Best Local Similarity 31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                             Query Match
Best Local Similarity
RESULT 516
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RESULT 523
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Best Local
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RESULT
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Length 217;

Length 217;

Length 217;

Length 217;

Length 217;

Length 217;

P P P P P P

RESULT 525

Best Loc RESULT 526

A D M E D

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003104542-A1.
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Human secreted polypeptide PR01078, SEQ ID NO:124
                22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM76387 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124 US203082717-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                             DB 6;
                                                                                                                                                                                                                                                                                             Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO02445 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003040061-A1.
                                                                  ABO41343 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068695-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AB003360 standard, protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003036127-A1.
                                                                                                                                                                                                       ABO36158 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068703-A1.
                                                                                                                                                                                                                                                                                                                                           AB043687 standard; protein; 217 AA
Human PRO polypeptide #62.
US2003068732-A1.
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Best Local Similarity 31.6%;
RESULT 537
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31.6%;
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Best Local Similarity 31.6%;
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31.6%;
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31.6%;
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31.6%;
                                                                                                                                  (GETH ) GENENTECH INC.
ry Match 22.6%;
t Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                          Best Local Similarity RESULT 534
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                                                                                                                      10-APR-2003
                                                                                                                                                          Query Match
                 Query Match
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                                                                                                                                                                          Best
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                                  DB 6; Length 217;
                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 217;
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                                                                                                                                                                        DB 6; Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6; Length 217;
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABMO7033 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068699-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM21127 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068707-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM09473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073175-A1.
                                                                                                                                                                                                                       ABMI1608 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064447-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM16005 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064463-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073184-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6;
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10-ARP-2003.
(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 6;
ery Match 22.6%; Pred. No. 4.1e-19;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 532
                              22.6%; Score 266.5; DB 6
31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                        Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO27566 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064451-A1.
                                                                              AB024597 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003065159-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US2003.0.03.03.4PR-2003.
(GETH ) GENENTECH INC.
Warch 22.6%; Sr
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31.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 531
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                           03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULY 533
ID ABM09473 stand
DE Human Ser
PN USPON
                 GENENTECH INC.
                                                                                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 527
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                                                  Best Local Similarity
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 03-APR-2003
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                                   Query Match
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RESULT 528
ID ABM160
DE Human
PN US2003
PD 03-APR

RESULT

P P P P P

Query

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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                       RESULT 553
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 6; Length 217;
                                    22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                             Length 217;
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20-MAR-2003.
(GETH ) GENENTECH INC.
ery Match 22.6%; Score 266.5; DB 6;
ery Match 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                           22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR87871 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
· US2003068718-A1.
                                                                                                                                                                                                                                                                                                                                                                      ABR94361 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
052003049177A1.
06-MR2-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR75868 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US200304429-A1.
06-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR71244 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003059880-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR93141 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064465-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR93446 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054478-A1.
                                                                                     ABR73684 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054468-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                            22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                 Human secreted/transmembrane protein (PRO) #62.
US2003054470-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22.6%; Score 266.5; DB 6 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 6
Pred. No. 4.1e-19;
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22.6%; Score 266.5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO27871 standard; protein; 217 AA
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                                                                                                                                     20-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                               Best Local Similarity
RESULT 544
ID ABR73684 standard; p1
DE Human secreted polype
PN US2003054468-A1.
PD 20-WAR-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 549
                                                                                                                                                                                          Best Local Similarity
RESULT 545
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RESULT 551
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RESULT 547
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Best Local Similarity
RESULT 552
US2003036130-A1
20-FEB-2003.
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                                    Query Match
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Length 217;
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                                                                                               Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                          DB 6; Length 217;
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                                                                                                                                                                                                                                                                                                           ABM04903 standard; protein; 217 AA.
Muman secreted polypeptide PRO1078, SEQ ID NO:124
US200306727-A1.
10-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM08863 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO:124
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 6;
22.7 Match 22.6%; Score 266.5; DB 6;
22.0%; Match 21.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003068714-A1.
ABO30006 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003064461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO39513 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003068776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM10388 standard; protein; 217 AA. Human secreted polypeptide PR01078, US2003069407-A1.
                                                                                                                                                        protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 217 AA
                                                                                                                                                                                          US20050cc.
10-APR-2003.
(GETH ) GENENTECH INC.
Watch 22.6%; SC
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S.
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PA (GETH ) GENENTECH INC.
QUERY MATCh 22.6%;
Best Local Similarity 31.6%;
RESULT 558
                                                       PD 03-APR-2003.

PA (GETH) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;
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Query Match
22.6%;
Best Local Similarity 31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO35548 standard; protein
Human PRO polypeptide #62.
US2003068758-Al.
                                                                                                                                                    ABO33215 standard; proteir
Human PRO polypeptide #62.
US2003068724-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                       Length 217;
                                                                                                                                                                                                                                                    DB 6; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                          DB 6; Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR86956 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM28142 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054476-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM15268 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068692-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                               ABR97168 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
                                                                                                        Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM10998 standard, protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049782-A1.
                                                                                                                                                                                                                                                    Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                            Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                    ABO23682 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032134-A1.
13-FEB-2003.
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Human secreted/transmembrane protein (PRO) #62.
US2003068733-A1.
                  ABO52059 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049768-A1.
                                                                                                                                                        ABOS2364 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049771-A1.
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13-MAR-2003.
(GETH ) GENENTECH INC.
"MATCh "...tity 31.6%;
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31.6%;
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2003.

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21H ) GENENTECH IN

SET LOCAL SIMILARITY 3

RESULT 563

ID ABO52364 standa-

DE Human PRO r.

PD 13-70030.

PD 13-7009
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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RESULT 566
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RESULT 564
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US2003054481-A1.
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RESULT 570
RESULT 562
1D ABO520
DE Human
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PD 13-MAR
PA (GETH
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13-FMATCH J GENERITECH INC.
ry Match
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try Match
try 31.6%; Score 266.5; DB 6; Length 217;
try Theral Similarity 31.6%; Pred. No. 4.1e-19;
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, (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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22.6%; Score 266.5; DB 6; Length 217;
t Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                             Length 217;
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ry Match 22.6%; Score 266.5; DB 6; Length 217;
t Local Similarity 31.6%; Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
12.6%; SCORE 266.5; DB 6; Length 217;
TOCAT Similarity 31.6%; Pred. No. 4.1e-19;
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068709-A1.
10-APR-2003.
                                                                                                                                                         ABM04234 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068716-A1.
                                                                                                                                                                                                                                                                                                                          Human secreted polypeptide PRO1078, SEQ ID.NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM33143 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003087374-A1.
08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM35380 standard; protein; 217 AA.
Human secreted polypeptide PRO1078; SEQ ID NO:124
US2003073179-A1.
                                                                                                                                                                                                                                                                         Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM07643 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068751-A1.
                                                                                          Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABC40733 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068684-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO50229 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049777-A1.
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Human secreted/transmembrane protein (PRO) #62.
US2003040055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein; 217 AA
                                            10-APR-2003.
(GETH ) GENENTECH INC.
22.6%;
rrv Match 22.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 573
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PA (GETH ) GENENTECH INC.

QUETY MATCh 22.6%;

Best Local Similarity 31.6%;
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US2003049773-A1.
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RESULT 578
                                                                                                             Best Local Similarity RESULT 572
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ABO41953 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049745-A1.
                                              PD 13-MAR-2003.

PA (GETH ) GENENTECH INC.

QUETY MATCh 22.6%;

BEST LOCAL Similarity 31.6%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                          Best Local Similarity RESULT 592
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RESULT 596
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                  Length 217;
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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 589
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Best Local Similarity 31.6%; Score 266.5; DB 6; Len RESULT 586

ID ABMO1184 standard; protein; 217 AA.

By US2003049770-A1.

PD 13-MAR-200?
                                                                                                                                                                                                                                                                                         ABM18445 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054480-A1.
               Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                             Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                              Score 266.5; DB 6;
Pred. No. 4.1e-19;
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A (GETH ) GENENTECH INC.

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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                       ABR97473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003059885-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR80573 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049740-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003073169-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM13438 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064457-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068711-A1.
                                                            ABO04275 standard; protein; 217 AA.
Human secreted(transmembrane protein (PRO) #62.
20.FEB-2003.
                                                                                                                                                                           ABO05905 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
21-FEB-2003.
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13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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               22.6%;
31.6%;
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
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     Query Match
Best Local Similarity
RESULT 581
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Best Local Similarity
RESULT 583
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                                                                                                                                           Best Local Similarity
RESULT 582
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RESULT 590
27-FEB-2003
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1/-AFK 2005.
(GETH) GENENTECH INC.
22.6%; Score 266.5; DB 6; Length 217;
FT T.ocal Similarity 31.6%; Pred. No. 4.1e-19;
Length 217;
                                                                                                                                                        Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6; Length 217;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QUELY, MACLO.
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 594
ID ABM32838 standard; protein; 217 AA.
DE Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                         ABMIDO83 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003067478-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM02404 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059886-A1.
DB 6;
                                                                                                                                                            DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM22652 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003087373-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM74863 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096353-A1.
22-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ÄBR96253 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
US2003054458-A1.
20-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                       USZUSZYZO

10-APR-2003.

(GETH ) GENENTECH INC.

22.6%; Score 266.5; DB 6;

RETY MATCh ...inlarity 31.6%; Pred. No. 4.1e-19;
Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                  RESULT 593
ID ABO38598 standard; protein; 217 AA.
DE Human secreted/transmembrane protein (PRO) #62.
PN US2003068773-A1.
                                                                             #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA79668 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003073173-A1.
                                                    ABO42563 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003049751-A1.
                                                                                         US2003042.
13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.6%;
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31.6%;
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 6; Length 217;
                                                      22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                              DB 6; Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 6; Length 217;
                                                                                                                                                                                                                              ABR86651 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM29667 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064456-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM23872 standard; protein; 217 AA.
Human secréted polypeptide PRO1078, SEQ ID NO:124
US2003068735-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM23262 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068753-A1.
                                                                                                   ABR86346 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049758-A1.
                                                                                                                                                                                                                                                                                                                                                            ABM16615 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064448-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM22042 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068742-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 22.6%; Score 266.5; DB 6. Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 606
                                                                                                                                                                                                                                                                                                            22.6%; Score 266.5; DB 6
31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO29091 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068693-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO37683 standard; protein; 217 AA
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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Best Local Similarity 31.6%;
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Lry Match
Best Local Similarity -
RESULT 600
ID ABR86346 stand*
DE Human secre
PN US20030
                                                                                                                                                                 (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 608
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                                                                                                                                                                                                                                                                                                                         Best Local Similarity
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                                                                                                                                                                                             Best Local Similarity RESULT 601
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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Best Local S
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ID ABO290
DE Human
PN US2003
PD 10-APR
PA (GETH
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Score 266.5; DB 6; Length 217;. Pred. No. 4.1e-19;
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Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
ry Match 22.6%; Score 266.5; DB 6; Length 217;
t Local Similarity 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                            Length 217;
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                                                                                                                                                                                                                                                                                                                                      ABM66396 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068737-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM75778 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104547-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM34058 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR96558 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                          ABM28447 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003082715-A1.
                                                                                                                                                                                        ABM28752 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003082716-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM34363 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
HUS2003100061-A1.
29-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                              Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
Best Local Similarity 31.6%; Fred. No. 4.1e-19; RESULT 609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO20294 standard; protein; ż17 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003032125-A1.
13-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO21209 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054454-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AB022124 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054477-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US20031000.
29-MAY-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
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                                                                                                                              22.6%;
31.6%;
                                                                                                                                                                                                                                                                            22.6%;
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PA (GETH ) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;

RESULT 612
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(GETH ) GENENTECH INC.
ery Match 22.6%;
st Local Similarity 31.6%;
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                                                                                                                                             Best Local Similarity RESULT 610
                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 611
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                                                                                                         01-MAY-2003.
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; DB 8; Length 384; . 22;
                                                                                                                                                                                                                                                                   werry match 7.0%; Score 82; DB 8; Length 366; Best Local Similarity 21.5%; Pred. No. 21; RESULT 1067
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 7.0%; Score 82; DB 4; Length 610;
Best Local Similarity 26.3%; Pred. No. 44;
RESULT 1073
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 4; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vuery Match 7.0%; Score 82; DB 3; Length 610; Best Local Similarity 26.3%; Pred. No. 44; RESULT 1071
                                                                                                      DB 8; Length 366;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 82; DB 3; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.0%; Score 82; DB 4; Length 610; 26.3%; Pred. No. 44;
Nicotiana tomentosiformis GPA1 ortholog, SEQ ID 38.
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mammalian toxicological response marker protein #8 WO200136684-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-525-2001.
(GETH ) GENENTECH INC.
7.0%; Score 82;
                                                                                                  Query Match 7.0%; Score 82; Best Local Similarity 21.5%; Pred. No. RESULT 1066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.0%; Score 82; 21.5%; Pred. No.
                                                                                                                                                            ADH78760 standard; protein; 366 AA.
Tobacco GPA1 ortholog, SEQ ID 40.
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                           ADH78762 standard; protein; 384 AA.
Tobacco GPAl ortholog, SEQ ID 42.
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADH78764 standard; protein; 384 AA. Tobacco GPA1 ortholog, SEQ ID 44. WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAM93654 standard; protein; 610 AA.
Human polypeptide, SEQ ID NO: 3522.
EP1130094-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAU29069 standard; protein; 610 AA.
Human PRO polypeptide sequence #46.
WO200168848-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY66653 standard; protein; 610 AA Membrane-bound protein PRO872.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAY92515 standard; protein; 610 AA
                                                                                                                                                                                                                                        (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA. Query Match 7.0%; Sc
                                                                                                                                                                                                                                                                                                                                                                                                                         (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA.
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                                                             (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 7.0%;
Best Local Similarity 26.3%;
RESULT 1070
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(INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (INCY-) INCYTE PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 1068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 1072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU04356 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200020604-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9963088-A2.
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                                               08-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-APR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Query Match 7.0%; Score 82; DB 6; Length 610; Best Local Similarity 26.3%; Pred. No. 44; RESULT 1078
                                                                                                                      DB 4; Length 610;
44;
                                                                                                                                                                                                                                       Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                          7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6; Length 610;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 6; Length 610;
44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 610
                                                                                                                                                                                                                                                               Novel human secreted and transmembrane protein PRO872. US2003032127-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABUS9069 standard; protein; 610 AA.
Növel human secreted or transmembrane protein PRO872.
US2002132252-A1.
                                      AAB65176 standard; protein; 610 AA.
Human PRO872 (UNQ439) protein sequence SEQ ID NO:113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABRÉGI82 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027278-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR65572 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036159-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU99512 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003040070-A1.
27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                            ABU84308 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 82; DB 6;
Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABU82581 standard; protein; 610 AA.
Human secreted/transmembrane protein PRO872.
US2003032023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB
44;
Best Local Similarity 26.3%; Pred. No. 44; RESULT 1074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 82;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.0%; Score 82; 26.3%; Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                        Score 82;
Pred. No.
                                                                                                                                                                      ABUS8445 standard; protein; 610 AA.
Human PRO polypeptide #46.
US2003027272-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU57991 standard; protein; 610 AA.
Human PRO polypeptide #23.
US2003027163-A1.
                                                                                                                                                                                                                                         7.0%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
                                                                                                      (GETH ) GENENTECH INC.
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                                                                                                                                  Best Local Similarity RESULT 1075
                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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RESULT 1080
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Best Local Similarity
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                                                                     WO200073454-A1.
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Length 217;
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22.6%; Score 266.5; DB 6; Length 217;
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PD 20-MAR-2003.

PA (GETH) GENENTECH INC.

Query Match

22.6%; Score 266.5; DB 6;

Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                     U3-WWR-2003.
(GETH ) GENENTECH INC.
(GETY Match 22.6%; Score 266.5; DB 6;
ery Match 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003068679-Al. 10-APR-2003. (GETH ) GENENTECH INC.
                                                                                                 ABR85736 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049753-A1.
13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                           22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                               Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM23567 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID.NO:124.
US2003068736-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM00269 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073172-A1.
                                                                                                                                                                                                                                                                                                                                                                                  ABM00574 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
117-APR-2003.
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AB029701 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003068700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO45593 standard, protein; 217 AA.
Human PRO polypeptide #62.
US2003073182-A1.
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Best Local Similarity 31.6%;
RESULT 627
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(GETH ) GENENTECH INC.
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RESULT 620
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Best Local Similarity
RESULT 621
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Best Local Similarity
RESULT 622
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05-UUN-2003.
(GETH ) GENENTECH INC.
lery Match 22.6%; Score 266.5; DB 6; Length 217;
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(GETH) GENENTECH INC.

Query Match 22.6%; Score 266.5; DB 6; Length 217;

Best Local Similarity 31.6%; Fred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 217;
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PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 636
                                                                                                                                                                                                                                                                                     Length 217;
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Pred. No. 4.1e-19;
                                      ABM20517 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003104557-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR81488 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049744-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR89701 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                         DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 6;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR92531 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064446-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM77912 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049783-A1.
                                                                                                                                                                                                                                     Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 628
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Pred. No. 4.1e-19;
                                                                                                                                                                                        ADA81395 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003092121-A1.
                                                                                                                                                                                                                                                                                                                                         ABO16631 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003027276-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO18257 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044920-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO22989 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003054461-Al.
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13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
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31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
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31.6%;
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13-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
Best Local Similarity
RESULT 647
                                                                                                                                        Best Local Similarity RESULT 648
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                                   22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted polypeptide PRO1078, SEQ ID NO:124. US203068702-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM03929 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068734-A1.
                                                                                                                                                           22.6%; Score 266.5; DB 6; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                            ABM13743 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064458-A1.
                                                                    Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003022121-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 6 Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane protein (PRO) #62. US20034460-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO30311 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064464-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO37073 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO41648 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068729-A1.
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10-ann
                                                                                                                                                                                                                                                                                                                                                                                                                                          22.6%;
31.6%;
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RESULT 642
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PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 31.6%;
RESULT 643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PD 10-APR-2003.

PA (GETH ) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                        03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                03-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                       Best Local Similarity
   US2003073171-A1.
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                                       Query Match
                                                                                                                                                           Query Match
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Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 654
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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 656
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22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                     22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                       Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GETH ) GENENTECH INC.

ry Match 22.6%; Score 266.5; DB 6;

t Local Similarity 31.6%; Pred. No. 4.1e-19;
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ID ABM16920 standard; protein; 217 AA.
DE Human secreted polypeptide PRO1078, SEQ ID NO:124
PN US2003040078-A1.
                                                                                                                                                                                                                                                                                                        13-MAR-2003.
(GETH ) GENENTECH INC.
(ery Match 22.6%; Score 266.5; DB 6;
ery Match 31.6%; Pred. No. 4.1e-19;
                                                                   ABMX5092 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104540-A1.
05-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                      ABG47789 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049747-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein (PRO) #62.
US20039779-A1.
                                                                                                                                                                                                                                            ABO47484 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049742-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO48399 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049750-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO51449 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO51754 standard; protein; 217 AA Human PRO polypeptide #62.
US2003049767-Al.
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Query Match
22.6%;
Best Local Similarity 31.6%;
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DB 6; Length 217;

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Score 266.5; DB 6; Length 217;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                 ABM75473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                            ABM25397 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104541-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM19907 standard, protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR71549 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003032133-A1.
                                                                                                                                                                                                                                                    05-JUN-2003.
(GETH ) GENENTECH.INC.
22.6%; Score 266.5; DB 6;
2ry Match ....ilarity 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                       ABO39208 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US203068775-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA81193 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049752-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO46813 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049762-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO47118 standard; protein; 217 AA
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                                                                                        US2003000...
10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S.
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05-JUN-2003.
(GETH ) GENENTECH INC.
22.6%; S
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13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; F
     22.6%;
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ry Match 22.6%;
r Local Similarity 31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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US2003049765-A1.
     Query Match
Best Local Similarity
RESULT 666
                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 668
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Best Local Similarity
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RESULT 667
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RESULT 669
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                                                                                                                                                                                                                                                                                                                                              Length 217;
                                                                           Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM16310 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064449-A1.
                                                                                                                                                                                                                                                ABR96863 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054462-A1.
                                                                                                                                                                                                                                                                                                                                                                                              ABM12218 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064445-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM24177 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064441-A1.
03-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM14658 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068696-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABM04519 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068712-A1.
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Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068730-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM09168 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073174-A1.
                                                                                                                                                                                                                                                                                                                                            DB 6;
                                                                                                                                                                                                 DB 6;
                                                                       Score 266.5; DB 6;
Pred. No. 4.1e-19;
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(GETH ) GENENTECH INC.
(AETY Match ....larity 31.6%; Score 266.5; DB 6;
                                                                                                                                                                                                 Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
ABO17952 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
                                                                                                                                        Human secreted/transmembrane protein (PRO) #62.
US2003032132-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                         ABO20904 standard; protein; 217 AA
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03-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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10-APR-2003.
(GETH ) GENENTECH INC.
worch 22.6%; S
                                                                       22.6%;
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03-APR-2003.
(GETH ) GENENTECH INC.
Match 'Trity 31.6%;
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31.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 664
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(GETH) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
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                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 658
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Best Local Similarity
RESULT 659
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RESULT 660
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Best Local Similarity
                                                                    Ouery Match
Best Local Similarity
RESULT 657
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Best Local Similarity
RESULT 662
                                    US2003044918-A1.
                                                     06-MAR-2003.
                                                                                                                                                                                13-FEB-2003.
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RESULT 661

DB 6; Length 217;

DB 6; Length 217;

6; Length 217;

RESULT 665

BESE

Length 217;

9

38

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                          Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                            22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                              22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                           ABR73379 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054467-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 6;
Pred. No. 4.1e-19;
Human secreted polypeptide PRO1078, SEQ ID NO:124 US2003036129-A1.
                                                                                                                                                                                                                         ABR84821 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040057-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR73074 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003027270-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM18140 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054469-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR76473 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044932-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                    Human secreted/transmembrane protein (PRO) #62.
US2003040053-A1.
27-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AB020599 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US20030312126-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PRO polypeptide #62.
US2003054466-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO25342 standard; protein; 217 AA. Human PRO polypeptide #62. US2003054463-A1.
                                                                                                        protein; 217 AA
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20-MAR-2003.
(GETH ) GENENTECH INC.
42.ch 22.6%; Sr
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31.6%;
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                                                       22.6%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                           (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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Best Local Similarity
RESULT 683
                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                   Best Local Similarity RESULT 676
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                                                                                                                                                                                          Local Similarity
                                                                                                        ABO06868 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .06-MAR-2003.
                                                                                                                                                                                                                                                                            27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-FEB-2003.
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                                                       Query Match
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03-APR-2003.
A (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 6; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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Query Match
22.6%; Score 266.5; DB 6; Length 217;
Beet Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 691
                                                                                               6; Length 217;
                                                                                                                                                                                                                               Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22.6%; Score 266.5; DB 6; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 217;
ABR94056 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059879-A1.
                                                                                                                                                                                                                                                                      Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM29972 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM05508 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003045700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM27227 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068760-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM15573 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                    ABR79963 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049738-A1:
                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                              Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Accounts standard; protein; 217 AA.

Human secreted/transmembrane protein (PRO) #62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO30921 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064468-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                       AB032910 standard, protein; 217 AA.
Human PRO polypeptide #62.
US2003064453-A1.
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
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06-MAR-2003.
(GETH ) GENENTECH INC.
22.6%;
                                         US2005.2.
27-MAR-2003.
(GETH ) GENENTECH INC.
22.6%;
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03-APR-2003.
(GETH ) GENENTECH INC.
"atch "laticy 31.6%;
                                                                                                                                                                                                                               22.6%;
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PA (GETH ) GENENTECH INC.
Query March 22.6%;
Best Local Similarity 31.6%;
RESULT 690
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                                                                                               Query Match
Best Local Similarity
RESULT 686
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                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                      RESULT 687
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Length 217;

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Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 6; Length 217; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003044924-Al. OG-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                       ABM26922 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068763-A1.
                                                                                                                                                                                                                                              ABR88176 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068720-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                    Score 266.5; DB 6;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                      ADA78488 standard; prótein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003073181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted/transmembrane protein (PRO) #62.
US2001049780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human secreted/transmembrane protein (PRO) #62. US2003036126-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO39818 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003068689-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO49924 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62
US2003049776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO05295 standard; protein; 217 AA
(GETH ) GENENTECH INC. 22.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US200302.
13-MAR-2003.
(GETH ) GENENTECH INC.
22.6%;
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
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RESULT 708
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
                                                                                                                                                           17-APR-2003.

1 (GETH) GENENTECH INC.

Query Match 22.6%;

Best Local Similarity 31.6%;
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1 (GETH) GENENTECH INC.

Query Match 22 6%;

Best Local Similarity 31.6%;
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Query Match
Best Local Similarity 31.6%;
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31.6%;
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(GETH ) GENENTECH INC.
                                                               Best Local Similarity RESULT 704
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Best Local Similarity
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                                                   Query Match
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                                           10-APR-2003.
(GETH ) GENENTECH INC.
ery Match 22.6%; Score 266.5; DB 6; Length 217;
ery Match 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                Length 217;
                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6; Length 217;
Pred, No. 4.1e-19;
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PA (G-JM-2003)

PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;

ID ABO49314 standard; protein; 217 AA.

PN US2003049774-A1.

PD 13-MAR-200.
                                                                                                                        ADMUNESS Standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068688-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 699
                                                                                                                                                                                                                                                                                 ABO42258 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049748-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                        ABO37988 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 6
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 6
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22.6%; Score 266.5; DB 6 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB20236 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003082767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO49619 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049775-A1.
13-PAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #62.
US2003049754-A1.
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; SC
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13-MAR-2003.
(GETH ) GENENTECH INC.
Match 'Trity 31.6%;
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PA (GETH) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
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Best Local Similarity
RESULT 696
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Best Local Similarity
RESULT 697
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                                                                                                  Best Local Similarity
RESULT 695
                                US2003068698-A1.
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Length 217;

Length 217;

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22.6%;
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(GETH ) GENENTECH INC.
22.6%;
St Local Similarity 31.6%;
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ry Match 22.6%;
t Local Similarity 31.6%;
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Best Local Similarity 31.6%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUN-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
               Best Local Similarity RESULT 723
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Best Local Similarity
                                                                                                                                            Best Local Similarity
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       Query Match
                                                                                                                                                                                                                                                                                     RESULT 725
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RESULT 730
                                                                              Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                        Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                   DB 7; Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR86041 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049759-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM10693 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064455-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM76692 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003054465-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR89396 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003073170-A1.
                                                                                                                                                                                                                                   ABR95886 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040073-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 7;
RESULT 713

ID ABR77078 standard; protein; 217 AA.

ID ABR77078 standard; protein; 217 AA.

DE Human secreted polypeptide PRO1078, SEQ ID NO:124

PN US2003044927-A1.
                                                                                                                          ABM17835 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040072-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 7;
Pred. No. 4.1e-19;
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20-0003.
(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 7;
ery Match 22.6%; Score 266.5; DB 7;
ery Match 31.6%; Pred. No. 4.1e-19;
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USACCE.

(GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

22.6%; Score 266.5; DB 7

(ery Match Cimilarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                   Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                              ABO21819 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054475-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted/transmembrane protein (PRO) #62. US2003032124-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted/transmembrane protein (PRO) #62
US2003064467-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO24292 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO19989 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                           US200302.
20-MAR-2003.
(GETH ) GENENTECH INC.
Warch 22.6%; SC
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03-APR-2003.
(GETH ) GENENTECH INC.
7-1 Match .'1arity 31.6%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%;
RESULT 720
                                                                                                                                                                                                                                                                                                   22.6%;
31.6%;
                                                                                 22.6%;
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31.6%;
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31.6%;
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Best Local Similarity
RESULT 722
ID ABR89396 stand
DE Human ser
PN USP
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Best Local Similarity
RESULT 719
                                                                                                                                                                                       Ouery Match
Best Local Similarity,
RESULT 715
                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 717
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Best Local Similarity
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Best Local Similarity
                                                                               Query Match
Best Local Similarity
RESULT 714
                                                                                                                                                                           27-FEB-2003.
                                                                                                                                                                                                                                                                                 27-FEB-2003.
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                                                                                                                                                                                                                                                                                                                              RESULT 716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT
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Lucir match 22.6%; Score 266.5; DB 7; Length 217; Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 729
                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ) 10-APR-2003.

4 (GETH) GENENTECH INC.

Query Match 22.6%; Score 266.5; DB 7; Length 217;

Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                  17.Akr-zvvv.
(GETH ) GENENTECH INC.
iry Match
22.6%; Score 266.5; DB 7; Length 217;
ir Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADMINISTRATE STANDARD; protein; 217 AA. Human secreted polypeptide PRO1078, SEQ ID NO:124 OF THE PROFESSO-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            אסר 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
082003104551-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003027273-A1.
                                                                    ABM12523 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073176-A1.
                                                                                                                                                                                                                                                      ABM05813 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM03014 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068764-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  אאר 17 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049757-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO34938 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003068728-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO46508 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2003049761-A1.
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03-APR-2003.
(GETH ) GENENTECH INC.
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                                                                                    Length 217;
                                                                                                                                                                                                                                                                                                                                               Length 217;
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(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 7; Length 217;
"F. Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                         Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 22.6%; Score 266.5; DB 7; Length 217; Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Luca polypeptide PRO1078, SEQ ID NQ:12

PD 27-MAR-2003.

PA (GETH) GENENTECH INC.

Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;

RESULT 738

ID ABM02099 standard; protein; 217 AA.

PN US2003059884-A1.

PD 27-MAR-2007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM02099 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059884-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM30582 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003064443-A1.
              ABR89091 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003015119-A1.
20-FEB-2003.
                                                                                                                                  ABR72464 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US200336120-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                        ABR74294 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003036161-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR80268 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003049739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM01489 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NQ:124
US2003059882-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted polypeptide PRO1078, SEQ ID NO:124. US2003068687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM12828 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073186-A1.
                                                                                                                                                                                                       22.6%; Score 266.5; DB 7; 31.6%; Pred. No. 4.1e-19;
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                                                                                  DB 7;
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                                                                                                                                                                                                                                                                                                                                             DB 7;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 737
                                                                                  22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                             Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                     ABO18562 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044921-A1.
                                                                                                                                                                                                                                                                                        22.6%;
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Best Local Similarity
RESULT 741
ID ABM30582 str
DE Humar
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                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 734
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 735
                                                                                 Query Match
Best Local Similarity
RESULT 733
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Best Local Similarity
RESULT 736
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Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM14353 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068686-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM09778 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003073178-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM34668 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104538-A1.
05-JUN-2003.
                                                           ABM24482 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064444-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                  ABO29396 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068697-A1.
10-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 266.5; DB 7
Pred. No. 4.1e-19;
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US2003068710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO38903 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068774-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO51144 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049781-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             #62
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Human secreted/transmembrane protein (PRO)
10/22003036158-A1.
20-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                               ABO31226 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 31.6%;
RESULT 745
                                                                                                                                                                      22.6%;
                                                                                                                                                                                                                                                                                                                                            22.6%;
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PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
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(GETH) GENENTECH INC.

Query Match

Best Local Similarity 31.64;
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Query Match 22.6%;
Best Local Similarity 31.6%;
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Best Local Similarity 31.6%;
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A (GETH) GENENTECH INC.

2 (GETH) GENENTECH INC.

2 6 8;

Best Local Similarity 31.68;
                                                                                                                          03-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-APR-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
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Best Local Similarity
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03-APR-2003.
(GETH ) GENENTECH INC.
                                                                    Best Local Similarity RESULT 761
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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                                                                                                         ABO31836 standard;
       US2003064450-A1.
                                                                                                                                                                                                                                                                                              10-APR-2003.
                                                         Query Match
                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                           DB 7; Length 217;
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                                                                                                                                                                                                                           Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABMO1794 standard, protein, 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003059883-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR90006 standard; protein; 217 AA.
Human secreted polypeptide PRO1078; SEQ ID NO:124
US2003073177-A1.
                                                                                                                                                                                                                                                                          ABR78893 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003054456-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR93751 standard; protein; 217 AA.
Human secreted polypeptide PR01078, SEQ ID NO:124
US2003054457-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003049764-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM13133 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM27532 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003064442-A1.
                                                                                                                                                        ABR77683 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7;
                                                                                                                                                                                                                                                                                                                                               Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-MAR-2003.
(GETH ) GENENTECH INC.
rry Match 22.6%; Score 266.5; DB 7
rry Match 31.6%; Pred. No. 4.1e-19;
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22.6%; Score 266.5; DB 7

it Local Similarity 31.6%; Pred. No. 4.1e-19;
     31.6%; Pred. No. 4.1e-19;
                                                                                                         Score 266.5; DB 7
Pred. No. 4.1e-19;
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Pred, No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                              ABO21987 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054482-A1.
                            Human PRO polypeptide #62.
US2003056151-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003u5xx.
20-MAR-2003.
(GETH ) GENENTECH INC.
22.6%; S
                                                                                                         22.6%;
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31.6%;
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31.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 755
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 756
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RESULT 757
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Best Local Similarity RESULT 751
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Best Local Similarity
                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                             Best Local Similarity
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                                                                                                                                                                                          US2003040067-A1.
                                                                                       20-FEB-2003.
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                                                                                                           Query Match
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22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                    Length 217;
                                                                                                                                                                                                                                                                                  Length 217;
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A (GETH ) GENENTECH INC.

Query Match 22.6%; Score 266.5; DB 7; Length 217;

Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 217;
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                                                                                                                                                                                           Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                         ABM08253 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068754-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM74558 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003096351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM33753 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096358-A1.
                                                                                                                                                    Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104556-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 7;
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                    Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO40123 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068681-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABO48704 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049756-A1.
                                                                      Human secreted/transmembrane protein (PRO) #62.0S2003068731-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR72769 standard; protein; 217 AA.
Human secreted polypeptide PRO1078,
UVS2003036122-A1.
20-FEB-2003.
                                                          protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                            US2003000...
10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; Sr
                                                                                                 USZOCZ-
10-APR-2003.
(GETH ) GENENTECH INC.
Match ''reity 31.6%;
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22-MAY-2003.
(GETH ) GENENTECH INC.
22.6%;
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05-UND-2003.
(GETH) GENENTECH INC.
22.6%;
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22-MAY-2003.
(GETH ) GENENTECH INC.
22.6%;
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SEQ ID NO:124

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Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
                                                                     Score 266.5; DB 7; Length 217;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                              ABM21432 standard; protein; 217 AA. '
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068744-A1.
10-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADMINISTRATED STANDARD STANDARD SECRETED NO:124 UNMAIN SECRETED NO:1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM75168 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003104544-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM33448 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003096357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Human secreted/transmembrane protein (PRO) #62.
US2003068694-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AB036768 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)·#62.
US2003068715-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABO37378 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003068726-A1.
                                                                                                                                                   ABM21737 standard; protein; 217 AA. Human secreted polypeptide PRO1078, US2003068741-A1.
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US2003049760-Al.
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22-MAY-2003.
(GETH ) GENENTECH INC.
Warch 22.6%; Sr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USZUCZC.
10-APR-2003.
(GETH ) GENENTECH INC.
7-- Match -- 12-64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-4PR-2003.
(GETH ) GENENTECH INC.
--- Match --- 13-rity 31.6%;
PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
QUERY MATCh 22.6%;
Best Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Match 22.6%;
Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PA (CBTH.) GENENTECH INC.

Query Match 22.6%;
Best Local Similarity 31.6%;
RESULT 782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-AFK-ZUCJ.
(GETH ) GENENTECH INC.
127 MATCh 22.6%;
127 MATCh 31.6%;
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1 (GETH) GENENTECH INC.

Query Match
Best Local Similarity 31.6%;
                                                                                                                                                                                                                                       10-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                          PA (GETH ) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 7; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 771
                                                                                                                                                                                                                                                                                                                                                                                                           DB 7; Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 7; Length 217;
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                                                                                                                                                                                                                                                                                        ABR85126 standard; protein; 217 AA,
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003040065-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted polypeptide PRO1078, SEQ ID NO:124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM76997 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM30277 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124.
US2003068723-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM17530 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003044928-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 266.5; DB 7;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM22957 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068757-A1.
        31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                         22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 266.5; DB 7
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22.6%; Score 266.5; DB 7 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO15106 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003044919-A1.
                                                                                               #62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human secreted/transmembrane protein (PRO) #62. US2003040077-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO20176 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003064459-A1.
                                                            ABO15411 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO)
US2003036121-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO17241 standard; protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2005.0...
03-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22.6%;
31.6%;
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A ) GENENTECH 1.
CY Match
Best Local Similarity PRSULT 777 10 ABO28176 stander
DE Human secretion PN US2003°
PN US2003°
PA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 779
                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
RESULT 772
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 773
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RESULT 774
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 775
        Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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SEE

1 2 2 C

22268

Length 217;

DB 7;

Length 217;

Length 217;

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Best Local Similarity 31.6%; Pred. No. 4.1e-19; RESULT 798
                                                                                                                                                                   Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                       22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22.6%; Score 266.5; DB 7; Length 217; 31.6%; Pred. No. 4.1e-19;
                   Score 266.5; DB 7; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 217;
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Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068771-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABM32107 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068708-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM32412 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068713-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM01497 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068761-Al.
                                                                                                                                                                                                                           ABM31802 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068680-A1.
                                                                                                                                                                                                                                                                                                                                                        ABM31192 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2003068762-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                        ADA82559 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003049755-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADBBE867 standard; protein; 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003054472-A1.
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(GETH.), GENENTECH INC.
22.6%; Score 266.5;
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Human secreted/transmembrane protein (PRO)
US2003087376-A1.
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10-APR-2003.
(GETH ) GENENTECH INC.
22.6%; S'
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                       22.6%;
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RESULT 795
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31.6%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
     (GETH ) GENENTECH INC.
                                                                                                                                                       GENENTECH INC
 PA (GETH ) concern Query Match Best Local Similarity RESULT 789
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
RESULT 797
                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best_Local Similarity RESULT 796
                                                                                                                                                                                        Best Local Similarity
RESULT 790
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Ob-NNV-zvus.
(GETH ) GENENTECH INC.
:ry Match 22.6%; Score 266.5; DB 7; Length 217;
:r f.ocal Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                        Length 217;
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ADG02592 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003207397-A1.
06-NOV-2003.
                                                                                                                                                                                                                                                                                           ADF95474 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003207398-A1.
                                                                                                                                           ADG01299 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2003207399-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PRO1078.
US2003073813-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG12289 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003207392-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADL32730 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003207396-A1.
06-NOV-2003.
                                                                                                                                                                                                                                        Score 266.5; DB 7;
Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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ry Match
22.6%; Score 266.5; DB 7;
t Local Similarity 31.6%; Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
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Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Aux/4261 standard, protein, 217 AA.
Human secreted/transmembrane protein (PRO) #62.
US2003211572-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADIGZ681 standard; protein; 217 AA.
Human apoptosis-associated protein SEQ ID 124.
WO2003058021-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE74873 standard; protein; 217 AA
                                                                                                                                                                                 US2003200.
06-NOV-2003.
(GETH) GENENTECH INC.
22.6%; SC
                                                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC. 22.6%; 21.6%; 31.6%;
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31.6%;
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Best Local Similarity 31.6%;
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US2003207395-A1.
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                                                                                                    Best Local Similarity
RESULT 799
                                                                                                                                                                                                                                                       Best Local Similarity RESULT 800
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RESULT 801
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RESULT 802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 806
ID ADE74261 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH08949 standard;
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(GETH ) GENENTECH INC.
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                                                    Length 217;
                                                                                                                                                                      Length 217;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 217;
                                                                                     ADVOOL human secreted and transmembrane protein PRO1078.
                                                                                                                                                                                                        ארס Standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
1052003215912-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH26054 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2003068770-A1.
                                                                                                                                                                                                                                                                                                                                                                             Novel human secreted and transmembrane protein PRO1078 US2003215911-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM31429 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2004048334-A1.
11-MAR-2004.
                                                  Score 266.5; DB 8;
Pred. No. 4.1e-19;
                                                                                                                                                                      22.6%; Score 266.5; DB 8; 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                       PD 20-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
22.6%; Score 266.5; DB 8;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                         0 20-NOV-2003.

(GETH ) GENENTECH INC.

Query Match 22.6%; Score 266.5; DB 8;

Best Local Similarity 31.6%; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 266.5; DB 8
Pred. No. 4.1e-19;
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PA (GETH) GENENTECH INC.
Query Match 22.6%; Score 266.5; DB 8
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
Human secreted/transmembrane protein (PRO) #62. US2003211574-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 266.5; DB 8
Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human PRO polypeptide #62.
                                                                                                                                                                                                                                                                                                                                                           ADG00517 standard; protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ54762 standard; protein; 217 AA.
Human PRO polypeptide #62.
US2004023321-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein; 217 AA
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... ) GENENTECH INC.
Best Local Similarity 31.6%; SC.
RESULT 815

ID ADJ64533 standard; prr.
DE Human PRO polyper
PN US20040383?-
PD 26-PP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Load; protein... polypeptide #62.
10-APR-2003.
PA (GETH) GENENTECH INC.
QUERY MAtch
Best Local Similarity 31.6%; Pre
RESULT 814
ID ADJ54762 standard; prov.
PN US2004023327.
PA (5-FFP)
PA (5-FFP)
                                                  22.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 10-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 22.6%;
Best Local Similarity 31.6%;
RESULT 813
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Best Local Similarity 31.6%;
RESULT 812
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(GETH ) GENENTECH INC.
                                                                                                                                                                             Best Local Similarity
RESULT 809
                                               Ouery Match
Best Local Similarity
RESULT 808
                                 13-NOV-2003.
                                                                                                                                                     20-NOV-2003.
                                                                                                                                                                        Query Match
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Length 217;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 22.6%; Score 266.5; DB 8; Length 217;
Best Local Similarity 31.6%; Pred. No. 4.1e-19;
RESULT 820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 217;
Score 266.5; DB 8; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                    Length 217;
                                                                                                                                                                                                                                                                                                     Score 266.5; DB 8; Length 217; Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 266.5; DB 8; Length 217; Pred. No. 4.1e-19;
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                                                        ADM36476 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US20040533S8-A1.
                                                                                                                                                                                                                                                                                                                                                        ADN37889 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078.
US2004091959-A1.
                                                                                                                                                                                                       ADM40281 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2004048335-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AED50119 standard; protein; 217 AA.
Novel human secreted and transmembrane protein PRO1078
US2005163766-A1.
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Pred. No. 4.1e-19;
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Best Local Similarity 31.6%; Pred. No. 4.1e-19;
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06-APR-2006.
(GETH ) GENENTECH INC.
22.6%; Score 266.5; DB 10;
                                                                                                                                                  Score 266.5; DB 8;
Pred. No. 4.1e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AEG62811 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
US2006073544-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEG72634 standard; protein; 217 AA.
Human secreted polypeptide PRO1078, SEQ ID NO:124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human placental protein; 217 AA. Human placental protein SEQ ID NO:1568. 17-NOV-2007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEG62199 standard; protein; 217 AA. Human secreted polypeptide PRO1078, US2006073545-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP24144 standard; protein; 217 AA. PRO polypeptide SEQ ID NO:1322. WO2004041170-A2.
                                                                                       US2004U5504.
18-MAR-2004.
(GETH ) GENENTECH INC.
22.6%; St.
 22.6%;
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31.6%;
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Local Similarity 31.6%;
                                                                                                                                                                                                                                                                 11-MAR-2004.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                              Best Local Similarity RESULT 818
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Query Match
Best Local Similarity
RESULT 817
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(NUVE-) NUVELO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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RESULT 835
                                                                                                                                                                                                                                         AEH17432 standard; protein; 217 AA.
Human tumor overexpressed cDNA protein product PRO1078 SEQ ID NO: 124.
                                                                                                                                                                                                                                                                                                                                                                       ADC32808 standard; protein; 244 AA.
Human novel contig-encoded polypeptide sequence, SEQ ID NO:2890.
WO2003029271-A2.
                                                                                                                                                                                        Length 217;
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Human T1 receptor-like ligand I protein SEQ ID NO:2.
US2005287606-A1.
                                                                                                      AEG88116 standard; protein; 217 AA.
Human PRO protein amino acid sequence - SEQ ID 124
US2006074227-A1.
                                                                                                                                                                                         DB 10;
                                                                                                                                                                                                                                                                                                                       22.6%; Score 266.5; DB 10; 31.6%; Pred. No. 4.1e-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22.3%; Score 262.5; DB 2; 31.1%; Pred, No. 1.1e-18;
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Pred. No. 8.4e-19;
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29-DEC-2005.
(HUMA-) HUMAN GENOME SCI INC.
22.3%; Score 262.5; DB 10
ery Match
22.3%; Score 262.5; DB 10
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Best Local Similarity 30.9%; Pred. No. 2.2e-18;
                                                                                                                                                                                        22.6%; Score 266.5; DB 1:31.6%; Pred. No. 4.1e-19;
                                                                         31.6%; Pred. No. 4.1e-19;
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Human p24 vesicle trafficking protein Hp24-1.
US2002132999-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW49099 standard; protein; 217 AA.
Human p24 vesicle trafficking protein Hp24-1.
WO9836068-A2.
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Human TI receptor-like ligand I.
WO9807881-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-JUN-2002.
(OXFO-) OXFORD BIOMEDICA UK LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-FEB-1998.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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31.6%;
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2002.
2002.
2002.
2ry Match
Best Local Similarity 3:
RESULT 831
ID AAW48334 stand?
DE Human T1 re
PN WO98079"
PD 26-7
Best Local Similarity 3
RESULT 826
ID AEG88116 standa-
DE Human PRO r
PN US2006r
PD 06-
                                                                                                                                                         06-APR-2006.
(GETH ) GENENTECH INC.
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                                                                                                                                                                                                   Best Local Similarity RESULT 827
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                                                                                                                                                                                                                                                                         US2006094864-A1.
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ADP04573 standard, protein, 208 AA.
Sea squirt protein with tissue specific expression in development Seq168.
JP2004057129-A.
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Human secreted protein sequence encoded by gene 18 SEQ ID NO:69.
WO200112775-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB74750 standard; protein; 201 AA.

Human secreted protein sequence encoded by gene 18 SEQ ID NO:59.
W02001775-A2.
22-FEB-2001.
(HUMA-) HUMAN GENOME SCI INC.
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Human secreted protein sequence encoded by gene 18 SEQ ID NO:68.
WO200112775-A2.
                                                                                                                                                                                                                                                                                                                    Score 236.5; DB 7; Length 201; Pred. No. 5.2e-16;
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(HUMA-) HUMAN GENOME SCI INC.
19.8%; Score 232.5; DB 4; Length 201;
                                                                                                                           20.1%; Score 236.5; DB 7; Length 201; 29.6%; Pred. No. 5.2e-16;
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Human protein highly expressed in cancer tissue SEQ ID NO 49.
WO2005076012-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-NOV-2004.
(FIVE-) FIVE PRIME THERAPEUTICS INC.
19.8%; Score 233.5; DB 8; Length 552;
Fr Local Similarity 30.8%; Pred. No. 4.6e-15;
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ry Match
t Local Similarity 28.7%; Pred. No. 9.9e-16;
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Pred. No. 9.9e-16;
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(KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
ADD44746 standard; protein; 201 AA.
Rat Protein Q63524, SEQ ID NO 10175.
W02003016475-A2.
27-FEB-2003.
                                                                                                                                                                                    ADD44750 standard; protein; 201 AA. Rat Protein Q63524, SEQ ID NO 10179-WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                  ABP69422 standard; protein; 208 AA.
Human polypeptide SEQ ID NO 1469.
WO200270539-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADU02684 standard, protein; 552 AA.
Novel human polypeptide seqid 1151.
WO2004093804-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-FEB-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                       20.1%;
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                                                       A (GEHO) GEN HOSPITAL CORP.
A (FARB) BAYER AG.
Query Match
                                                                                                                                                                                                                                                                             (GEHO ) GEN HOSPITAL CORP. (PARB ) BAYER AG.
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Best Local'Similarity
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                                                                                                                                                                                                                                                           27-FEB-2003
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15-JAN-2004.
(ROSE/) ROSEN C A.
(HASE/) HASELTINE W A.
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                                                                                                                                                                              BABARA
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Albumin fusion protein related therapeutic protein X, SEQ ID No 2073.
US2004010134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADL/8593 standard; protein; 201 AA.
Albumin fusion protein related therapeutic protein X, SEQ 1D No 2075.
US2004010134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ68911 standard; protein; 201 AA.
Human heat mitochondrial protein as a therapeutic target SeqID717.
WQ2003087768-A2.
                                                                                                                                                                                                  Length 201;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19.8%; Score 232.5; DB 7; Length 201; 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 8; Length 201;
                                                                                                                                                          (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABG65324 standard; protein; 201 AA.
Human albumin fusion protein #1999.
N0200177137-A1.
18-OCT-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
st Local Similarity 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19.8%; Score 232.5; DB 7; 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 232.5; DB 7;
Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                               DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 7;
                                                                                                                                                                                                                                                                                                                                                     MCZCT-2001.

(HU-OCT-2001.

(HU-OCT-
29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 232.5; DB 7
Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19.8%; Score 232.5; DB 8 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-SEP-2003.
(CHIL-) CHILDREN'S HOSPITAL MEDICAL CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD44752 standard; protein; 201 AA.
Human Protein Q15363, SEQ ID NO 10181.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADD44748 standard; protein; 201 AA.
Human Protein Q15363, SEQ ID NO 10177.
27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP65317 standard; protein; 201 AA.
Human coated vesicle membrane protein.
WO2003072827-A1.
                                                   ABG65327 standard; protein; 201 AA. Human albumin fusion protein #2002 WO200177137-A1.
                                                                                                                                                                                                                                                                             ABG65326 standard; protein; 201 AA. Human albumin fusion protein #2001.WO200177137-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-FEB-2003.
(GEHO ) GEN HOSPITAL CORP.
(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MITO-) MITOKOR.
(BUCK-) BUCK INST AGE RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ROSE/) ROSEN C A. (HASE/) HASELTINE W A.
                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 845
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
RESULT 846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 847
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
RESULT 851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 852
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Best Local Similarity
RESULT 849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-2003
                                                                                                                                            18-OCT-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match
                               RESULT 844
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ABM81493 standard; protein; 201 AA.
Tumour-associated antigenic target (TAT) polypeptide PRO4793, SEQ:3848.
WO2004030615-A2.
                                            ADL/8594 standard; protein; 201 AA.
Albumin fusion protein related therapeutic protein X, SEQ ID No 2076.
US2004010134-A1.
15-JAN-2004.
(ROSE/) ROSEN C A.
(HASE/) HASELTINE W A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-APR.2006.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
19.8%; Score 232.5; DB 10; Length 201;
ery Match
29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.
2ry Match
3t Local Similarity 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 10; Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                           15-APR-2004.
(GETH ) GENENTECH INC.
(ery Match 19.8%; Score 232.5; DB 8; Length 201;
19.8%; Score 232.5; DB 8; Length 201; 29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                          Score 232.5; DB 8; Length 201; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 201;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADK62498 standard; protein; 215 AA.
Disease treating protein complex-derived protein #375.
                                                                                                                                                                                                                                                                                                                                Score 232.5; DB 8;
Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 8;
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9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19.8%; Score 232.5; DB 10.29.2%; Pred. No. 1.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 229.5; DB 6
Pred. No. 3.1e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADP54500 standard; protein; 201 AA.
Human PRO protein sequence SEQ ID NO:476.
WO2004039956-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEH08866 standard; protein; 201 AA.
Therapeutic protein HCFCU69, SEQ ID 2075.
US2006084794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AEH08864 standard; protein; 201 AA.
Therapeutic protein HACAB58, SEQ ID 2073.
US2006084794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEH08867 standard; protein; 201 AA.
Therapeutic protein HCB3Z39, SEQ ID 2076.
US2006084794-A1.
                                                                                                                                                                                                                              ADN04050 standard; protein; 201 AA. Antipsoriatic protein sequence #220 WO2004028479-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR52923 standard, protein, 215 AA.
Protein sequence #SEQ ID 711.
EP1258494-Al.
                                                                                                                                                                                                                                                                    WO2004uzc..
08-APR-2004.
(GETH ) GENENTECH INC.
19.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19.5%;
                                                                                                                                                                      Query Match
Best Local Similarity 29.2%;
RESULT 854
              Best Local Similarity
RESULT 853
                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
RESULT 855
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Best Local Similarity
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Length 207;

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/ Match 16.4%; Score 193.5; DB 2; Length 163; Local Similarity 36.7%; Pred. No. 1.3e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 181.5; DB 4; Length 203;
Pred. No. 3.1e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AEB88199 standard; protein; 158 AA.
Human protein highly expressed in cancer tissue SEQ ID NO 50.
WO2005076012-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-AUG-2005.
(FIVE-) FIVE PRIME THERAPEUTICS INC.
ery Match 14.1%; Score 165.5; DB 9; Length 158;
ery Match 29.1%; Pred. No. 1.1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 163.5; DB 7; Length 135;
Pred. No. 1.4e-08;
                                                                                                                                                                                                                                                                          PD 10-JAN-2001.
PA (HELI-) HELIX RES INST.
Query Match 15.7%; Score 185; DB 4; Length 146;
Best Local Similarity 34.6%; Pred. No. 8.4e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE10062 standard; protein; 254 AA.
Novel protein-related contig polypeptide seguence #650
WO2003054152-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Abbe4154 Standard; protein; 203 AA.
Drosophila melanogaster polypeptide SEQ ID NO 19854.
WO200111042-A2.
                                                                                                                                                                                                                                    AAB88445 standard; protein; 146 AA.
Human membrane or secretory protein clone PSEC0228
EP1067182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC31097 standard; protein; 135 AA.
Human novel polypeptide sequence, SEQ ID NO:1179-
WO2003029271-A2.
                       vuery Match
Best Local Similarity 28.2%; Pred. No. 2.9e-12;
RESULT 871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15.7%; Score 185; DB 9; 34.6%; Pred. No. 8.4e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 166; DB 7;
Pred. No. 1.8e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB24228 standard; protein; 240 AA.
Human vesicle associated protein 7 SEQ ID NO:7.
WO200060082-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 166; DB 4;
Pred. No. 1.8e-08
                                                                              AAW70458 standard, protein, 163 AA.
Human T1-receptor ligand III splice variant 1.
WO9838311-A1.
                                                                                                                                                                                                                                                                                                                                                                                                ADY63253 standard; protein; 146 AA.
Human clone PSEC0228 protein, SEQ ID 258
EP1514933-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAU31151 standard; protein; 254 AA.
Novel human secreted protein #1642.
WO200179449-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                               03-SEP-1998.
(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14.1%; 27.1%;
               (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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Best Local Similarity
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(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-JUL-2003.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-OCT-2001.
(HYSE-) HYSEQ INC.
22-JUN-2006
                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEJ50287 standard; protein; 207 AA.
Saccharomyces cerevisiae stress-related protein (SRP) - SEQ ID 206.
US2006137043-A1.
                                                                                                                                                                                                             DB 10; Length 203;
                                                                                                                                                                                                                                                                                                                                                              Score 218.5; DB 8; Length 201; Pred. No. 4e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 183;
                                                     19.5%; Score 229.5; DB 7; Length 215; 29.3%; Pred. No. 3.1e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17.2%; Score 202.5; DB 8; Length 188; 30.3%; Pred. No. 1.8e-12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADL31427 standard; protein; 188 AA.
Human protein encoded by a full length cDNA clone SeqID 3460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 210;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 208;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 207;
                                                                                                                                                                                                                                                                                                                                                                                                                    ABBS9997 standard; protein; 208 AA.
Drosophila melanogaster polypeptide SEQ ID NO 6783.
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB58697 standard; protein; 210 AA.
Drosophila melanogaster polypeptide SEQ ID NO 2883.
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 215.5; DB 5;
Pred. No. 7.3e-14;
                                                                                            Amnuvilz standard; protein; 203 AA.
Protein aggregation-related protein SEQ ID NO:6.
WO2006091964-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 217; DB 4;
Pred. No. 6.1e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 216; DB 4;
Pred. No. 7.8e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADT87228 standard; protein; 207 AA.
Yeast Stress-related protein from gene YOR016C.
WO2004092398-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17.1%; Score 201; DB 8; 28.2%; Pred. No. 2.9e-12;
                                                                                                                                                                                                           Score 221.5; DB Pred. No. 2e-14;
                                                                                                                                                                                                                                                          ADS10510 standard; protein; 201 AA.
Human therapeutic protein - SEQ ID 747.
WO2004080148-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABP69421 standard; protein; 183 AA. Human polypeptide SEQ ID NO 1468. WO200270539-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAM93624 standard; protein; 188 AA.
Human polypeptide, SEQ ID NO: 3460.
EP1130094-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-MAR-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18.4%;
26.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18.4%;
28.7%;
                                                                                                                                                                                                                                                                                                                                                              18.6%;
28.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                           18.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-SEP-2001.
(HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOO4.

LAS-) RES ASSOC B.

LY Match
Best Local Similarity 3
RESULT 869
ID ADT87228 stand?
DE Yeast Stree?
PN WO2004^C
PD 28-C
PA
                                                                                                                                                                                       (UYAL-) UNIV ALABAMA.
                 27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                      Best Local Similarity RESULT 863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                         23-SEP-2004.
(NUVE-) NUVELO INC.
                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 864
                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local Similarity
RESULT 865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 866
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 867
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-2002.
(HYSE-) HYSEQ INC.
EP1338608-A2.
                                                                                                                                                                       31-AUG-2006
                                                       Query Match
                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
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Length 146;

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Length 240;
                                                                                                                                                       Length 240;
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 240;
                                                                                                                                                                                                                                                                                         4; Length 240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 136;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 8; Length 225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 8; Length 225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 228;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila melanogaeter polypeptide SEQ ID NO 22431 WO200171042-A2.
                                                                                                           GC-APR-2001.
(ZYMO) ZYMOGENETICS INC.
(ZYMO) ZYMOGENETICS INC.
13.8%; Score 162.5; DB 4;
ery Match
' n'milarity 22.2%; Pred. No. 4e-08;
                 13.8%; Score 162.5; DB 3; 22.2%; Pred. No. 4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13.7%; Score 161.5; DB 8; 29.2%; Pred. No. 2.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                           13.8%; Score 162.5; DB 5; 22.2%; Pred. No. 4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADX66737 standard; protein; 225 AA.
Plant full length insert polypeptide segid 37580.
US2004034888-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADX89364 standard; protein; 225 AA.
Plant full length insert polypeptide seqid 52028
US2004034888-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13.7%; Score 161.5; DB 8 25.2%; Pred. No. 4.6e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 156.5; DB 8 Pred. No. 1.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4;
                                                                                                                                                                                                                                                                                       13.8%; Score 162.5; DB 22.2%; Pred. No. 4e-08;
                                                                                                                                                                                                  AAG81378 standard; protein; 240 AA.
Human AFP protein sequence SEQ ID NO:274.
WO200129221-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13.0%; Score 153;
                                                              AAB62393 standard; protein; 240 AA.
Human type I membrane protein ztsll
WO200129223-A2.
                                                                                                                                                                                                                                                                                                                                      ABP64862 standard; protein; 240 AA Human protein SEQ ID 522. WO200259260-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              228 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADK66189 standard; protein; 136 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABB65213 standard; protein;
(INCY-) INCYTE PHARM INC
                                                                                                                                                                                                                                                    26-APR-2001.
(ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HAYA/) HAYASHIZAKI Y.
(KAMI/) KAMIYA M.
(SUZU/) SUZUKI T.
(HIRO/) HIROKAWA H.
                                                                                                                                                                                                                              Best Local Similarity
RESULT 881
ID AAG81378 standard; pr.
DE Human AFP protein seq
PN WO200129221-A2.
PD 26-APR-2001.
PA (ZYMO) ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZHOU/) ZHOU Y.
KOVA/) KOVALIC D K.
SCRE/) SCREEN S E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human phX-6 protein.
US2004019006-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 883
             Query Match
Best Local Similarity
RESULT 880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TABA/) TABASKA J E. (CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TANA/) TANAKA T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAKAO K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LIUJ/) LIU J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LIU J.
                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ΩI
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ABM80706 standard; protein; 214 AA.
Tumour-associated antigenic target (TAT) polypeptide PRO19615, SEQ:1822.
WO2004030615-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADJ70121 standard; protein; 214 AA.
Human heat mitochondrial protein as a therapeutic target SeqID1927.
W02003087768-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 203;
Best Local Similarity 20.4%; Pred. No. 3.7e-07;
RESULT 887

DB AEL96281 standard; protein; 228 AA.

DE Drosophila melanogaster survival essential protein, SEQ:1173.

PN US713558-81.
                                                                                                                                                                                                                                                                                                                                                                Length 219;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 203,
                                                                                                                           Length 228;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 216;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 214;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Arabidopsis thaliana protein; 253 AA.

EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAG44244 standard; protein; 216 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 55397.
BP1031405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Auk62494 standard; protein; 203 AA.
Disease treating protein complex-derived protein #373.
EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ..
9
                                                                                                                                                                                                                                                                                                                                                                Score 146.5; DB 8;
Pred. No. 1.7e-06;
                                                                                                                                                                               ADX78578 standard; protein; 219 AA.
Plant full length insert polypeptide segid 47944.
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11.9%; Score 140.5; DB 7; 26.3%; Pred. No. 6.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Abovovi standard; protein; 227 AA.
Human protein having hydrophobic domain, HP03977.
WO200149728-A2.
                                                                                                                             13.0%; Score 153; DB 11; 20.4%; Pred. No. 3.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 140.5; DB 6
Pred. No. 6.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11.6%; Score 137; DB 8; 24.0%; Pred. No. 1.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11.5%; Score 135; DB 3; 23.7%; Pred. No. 2.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11.5%; Score 135; DB 3; 23.7%; Pred. No. 3.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 137; DB 7;
Pred. No. 1.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                       Protein sequence #SEQ ID 707.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-0CT-ZUUS.

(MITO-) MITOKOR.

(BUCK-) BUCK INST AGE RES.

11.6%; S
                                                                                                                                                                                                                                                                                                                                                                12.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11.9%;
26.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GETH ) GENENTECH INC.
                                                                                        14-NOV-2006.
(APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                   LIU J.
ZHOU Y.
KOVALIC D K.
SCREEN S E.
TABASKA J E.
                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-NOV-2002.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                                                              (CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-APR-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                  Query Match
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                                                                                                                                                                                                                                                                                                                             (TABA/)
                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 889
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RESULT 904
ID AAG103
DE Arabic
PN EP103
                                                  Lucay match 11.4%; Score 134.5; DB 4; Length 227; Best Local Similarity 25.1%; Pred. No. 3.2e-05; RESULT 896
                                                                                                                                                                                                                               11.4%; Score 134.5; DB 4; Length 227; 25.1%; Pred. No. 3.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                           11.4%; Score 134.5; DB 7; Length 227; 25.1%; Pred. No. 3.2e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vuery Match 11.3%; Score 132.5; DB 7; Length 212;
Best Local Similarity 23.5%; Pred. No. 4.7e-05;
RESULT 902
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 218;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11.0%; Score 130; DB 3; Length 208; 25.2%; Pred. No. 8.4e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 203;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADK62524 standard; protein; 212 AA.
Disease treating protein complex-derived protein #388.
EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAG10323 standard; protein; 208 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 8594.
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF69317 standard, protein; 218 AA.
Human lung specific protein sequence SEQ ID NO:74.
W22003102137-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADY22460 standard; protein; 143 AA.
Plant full length insert polypeptide segid 70244
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
11.3%; Score 133; DB 5;
Best Local Similarity 22.0%; Pred. No. 3.9e-05;
RESULT 899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 133; DB 5;
Pred. No. 3.9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match . 11.3%; Score 133; DB 8;
Best Local Similarity 24.2%; Pred. No. 4.3e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 131.5; DB 8
Pred. No. 3.4e-05;
                                                                                                                   AAU27676 standard; protein; 227 AA.
Human full-length polypeptide sequence #1.
WO200164834-A2.
                                                                                                                                                                                                                                                                                      ADD19280 standard; protein; 227 AA. Human secreted protein from gene 22. WQ2003052377-A2. 26-JUN-2003. (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADH48778 standard; protein; 203 AA. NOV26 protein sequence, SEQ ID 62. WO200268652-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALJ61737 standard; protein; 203 AA. Human NOV26 protein, SEQ ID NO: 62. EP1686175-A2.
12-JUL-2001.
(PROT-) PROTEGENE INC.
(SAGA) SAGAMI CHEM RES CENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11.2%;
30.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 22.0%;
RESULT 900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-FEB-2004.
(LIUD) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-AUG-2006.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-DEC-2003. (DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-SEP-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                              Best Local Similarity RESULT 897
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 903
                                                                                                                                                                                   07-SEP-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 898
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 901
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AEC64808 standard; protein; 219 AA.
Deregulated, liver cancer-associated protein, SEQ ID NO:123
W02005085861-A2.
(ORID-) ORIDIS BIOMED FORSCHUNGS & ENTWICKLUNGS.
                                                                                 Length 212;
                                                                                                                                                                                                                                                                                                                                                                                                                            10.6%; Score 125; DB 4; Length 237; 25.2%; Pred. No. 0.00034;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 219;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 10.5%; Score 124; DB 7; Length 219; Local Similarity 24.9%; Pred. No. 0.00039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.5%; Score 123; DB 7; Length 219; 24.6%; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 10.5%; Score 124; DB 9; Length 219; Local Similarity 24.6%; Pred. No. 0.00039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 10.5%; Score 123; DB 7; Length 193; Local Similarity 25.5%; Pred. No. 0.00041;
                                                                                                                                                                                                                                                     Score 126; DB 7; Length 225; Pred. No. 0.00025;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rice abiotic stress responsive polypeptide SEQ ID NO:5225. W02003008540-A2.
                                                                                                                                              ADK62522 standard; protein; 225 AA.
Disease treating protein complex-derived protein #387
EP1338608-A2.
AAG10322 standard; protein; 212 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 8593
                                                                                 Score 130; DB 3;
Pred. No. 8.6e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10.5%; Score 124; DB 7; 24.9%; Pred. No. 0.00039;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD48307 standard; protein; 219 AA.
Rat Protein CAA06212, SEQ ID NO 14005.
WO2003016475-A2.
27-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD48311 standard; protein; 219 AA.
Rat Protein CAA06212, SEQ ID NO 14009.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD48309 standard; protein; 219 AA.
Human Protein P49755, SEQ ID NO 14007.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Protein P49755, SEQ ID NO 1367. WO200316475-A2.
                                                                                                                                                                                                                                                                                                               AA013679 standard; protein; 237 AA. Human polypeptide SEQ ID NO 27571. W0200164835-A2. 077-SEP-2001. (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SYGN ) SYNGENTA PARTICIPATIONS AG.
                                                                                                                                                                                                                                                        10.7%;
20.3%;
                                                                                 Query Match 11.0%;
Best Local Similarity 25.2%;
RESULT 905
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-PEB-2003.
(GEHO ) GEN HOSPITAL CORP.
(FARB ) BAYER AG.
                                                                                                                                                                                                          27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 906
                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 907
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Best Local Similarity
                                         EP1033405-A2.
06-SEP-2000.
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Length 205;

Length 207;

Length 214;

Length 224;

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Score 119.5; DB 3; Length 204; Pred. No. 0.001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9.7%; Score 114.5; DB 4; Length 100; 40.4%; Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 114.5; DB 8; Length 119; Pred. No. 0.0016;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9.6%; Score 113.5; DB 7; Length 203; 23.8%; Pred. No. 0.0044;
                                                                                                                                                                                                                                          DB 3; Length 225;
                                                     AAG16692 standard, protein; 205 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 17436.
PP1031405-A2.
06-SEP-2000.
                                                                                                                                                                     AAG16691 standard; protein; 225 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 17435.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                      AAGO7344 standard; protein; 214 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 4462.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAG07343 standard; protein; 224 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 4461.
06-SEP-2000.
                                                                                                                       10.2%; Score 119.5; DB 3; 27.3%; Pred. No. 0.001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 119; DB 3;
Pred. No. 0.0013;
                                                                                                                                                                                                                                                                                                                                                                       10.1%; Score 119; DB 2; 20.6%; Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.1%; Score 119; DB 3; 20.5%; Pred. No. 0.0013;
                                                                                                                                                                                                                                        Score 119.5; DB
Pred. No. 0.0012;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABG20510 standard; protein; 100 AA.
Novel human diagnostic protein #20501.
WO200175067-A2.
                                                                                                                                                                                                                                                                                    protein; 207 AA. protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADK66184 standard; protein; 119 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADES5548 standard; protein; 203 AA.
Rat Protein Q63584, SEQ ID NO 1365.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADT60183 standard; protein; 244 AA. Plant polypeptide, SEQ ID 10260. US2004216190-A1.
        10.2%;
                                                                                                                                                                                                                                        10.2%;
27.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.1%;
20.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHO ) GEN HOSPITAL CORP. (FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HAYA/) HAYASHIZAKI Y.
(KAMI/) KAMIYA M.
(SUZU/) SUZUKI T.
(HIRO/) HIROKAWA H.
                                                                                                                                                                                                                                                                                                       H. contortus pHC-2 p
DE19704024-A1.
16-APR-1998.
(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse phX-6 protein.
US2004019006-A1.
29-JAN-2004.
   Query Match
Best Local Similarity
RESULT 922
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RESULT 924
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RESULT 928
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Best Local Similarity
RESULT 929
                                                                                                                                     Best Local Similarity RESULT 923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 926
                                                                                                                                                                                                                                                                                         AAW59051 standard;
                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 930
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NAKA/) NAKAO K.
(TANA/) TANAKA T.
                                                                                                                                                                                                                       06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-SEP-2000.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-OCT-2001
                                                                                                                       Query Match
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RESULT 925
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                                                                                                                                                                                                                                                                                         22262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human heat mitochondrial protein as a therapeutic target SeqID719, WO2003087768-A2.
                                                                                                                                                                                                                                                                                                                                                                                    PD 08-APR-2004.
PA (GETH) GENENTECH INC.
Query Match 10.5%; Score 123; DB 8; Length 219;
Best Local Similarity 24.6%; Pred. No. 0.00049;
RESULT 916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PD 13-MAY-2004.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.6%; Pred. No. 0.00049;
RESULT 917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 123; DB 10; Length 219;
Pred. No. 0.00049;
      Length 219;
                                                                                                                                                    10.5%; Score 123; DB 7; Length 219; 24.6%; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                     Score 123; DB 7; Length 219; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 219;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADY70643 standard; protein; 219 AA.
Human nicastrin/BACE1/Psen2-complex member, TMP21 protein.
WO2005023833-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAG16693 standard; protein; 204 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 17437.
EP1033405-A2.
06-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABP52942 standard; protein; 218 AA.
Human lung specific protein sequence SEQ ID NO:185.
WO200264788-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AEK18280 standard; protein; 219 AA.
Human transmembrane trafficking protein, TMP21.b.
US2006204503-A1.
     10.5%; Score 123; DB 7; 24.6%; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PD 21-MAY-2004.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.6%; Pred. No. 0.00049;
RESULT 918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10.5%; Score 123; DB 9; 24.6%; Pred. No. 0.00049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.3%; Score 121; DB 5; 24.6%; Pred. No. 0.00079;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO protein; 219 AA. Howan PRO protein sequence SEQ ID NO:125.
                                                  ADD48113 standard; protein; 219 AA.
Human Protein P49755, SEQ ID NO 14011.
WO2003016475-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP23066 standard; protein; 219 AA.
PRO polypeptide SEQ ID NO:160.
WO2004041170-A2.
                                                                                                                                                                                                       ADJ68913 standard; protein; 219 AA.
                                                                                                                                                                                                                                                                                                                                                    ADN03686 standard; protein; 219 AA.
                                                                                                                                                                                                                                                                                                                                                                     Antipsoriatic protein sequence #40. WO2004028479-A2.
                                                                                                                                                                                                                                            (MITO-) MITOKOR.

(BUCK-) BUCK INST AGE RES.

10.5%; Sc
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24.6%;
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(BIOJ ) BIOGEN IDEC MA INC.
                                                                                                27-FEB-2003.
(GEHO ) GEN HOSPITAL CORP.
(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -1,
-2005.
-2005.
-2Ty Match
Best Local Similarity 2.
RESULT 919
ID AEK18280 standar
DE Human transf
PN US200627
PD 14-7
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Query Match
Best Local Similarity
RESULT 913
                                                                                                                                                    Query Match
Best Local Similarity
RESULT 914
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Best Local Similarity
RESULT 915
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09-MAR. 2000.
(UYNY) UNIV NEW YORK STATE.
                                                                                                                                                                                                                                                                                                                   Match 8.6%;
Local Similarity 19.7%;
                                                                                                                                                                              Match 8.6%;
Local Similarity 19.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 8.2%;
Local Similarity 24.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-FEB-2001.
(HELI-) HELIX RES INST.
(TABA/) TABASKA J E. (CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 944
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                                        Query Match
Best Local Similarity
RESULT 940
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                       Best Loc
RESULT 942
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                                                                                                                                                                                                                                      82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            / Match 9.1%; Score 107; DB 8; Length 204; Local Similarity 19.8%; Pred. No. 0.021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8.8%; Score 103; DB 4; Length 427; 25.1%; Pred. No. 0.16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 211;
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                        Length 244;
                                                                                                                                                            Length 204;
                                                                                                                                                                                                                                                                                                                   Length 136;
                                                                                                                                                                                                              AEB88231 standard; protein; 136 AA.
Human protein highly expressed in cancer tissue SEQ ID NO
WO2005076012-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAG13954 standard, protein; 214 AA.

Arabidopsis thaliana protein fragment SEQ ID NO: 13637.

BEP1031405-A2.

Query Match

Best Local Similarity 19.7%; Pred. No. 0.06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG13953 standard; protein; 237 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 13636.
EP103405-A2.
06-SEP-2000
                                                                              AAG07345 standard, protein; 204 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 4463.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABB64424 standard; protein; 427 AA.
Drosophila melanogaster polypeptide SEQ ID NO 20064.
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                       ABP52926 standard; protein; 211 AA.
Human lung specific protein sequence SEQ ID NO:169.
WO200264788-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADJ63854 standard; protein; 204 AA.
Plant lipid metabolism protein Jb027 SEQ ID NO:56.
WO2004013304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADX95475 standard; protein; 102 AA.
Plant full length insert polypeptide segid 58139
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8.8%; Score 103; DB 3;
19.7%; Pred. No. 0.07;
                      9.3%; Score 110; DB 8; 27.7%; Pred. No. 0.013;
                                                                                                                                                        Score 109; DB 3;
Pred. No. 0.013;
                                                                                                                                                                                                                                                                    18-AUG-2005.
(RIVE-) FIVE PRIME THERAPEUTICS INC.
ery Match
9.2%; Score 108; DB 9;
ery Match
77.2%; Pred. No. 0.0093;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       9.2%; Score 108; DB 5; 23.8%; Pred. No. 0.018;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AMPA-) FUNDACAO AMPARO A PESQUISA DO ESTADO. ry Match tocal Similarity 22.1%; Pred. No. 0.024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADY64838 standard; protein; 206 AA.
S. mansoni protein SEQ ID 256.
WO2005023979-A2.
                                                                                                                                                          9.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BADI ) BASF PLANT SCI GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                               22-AUG-2002.
(DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
      (KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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Best Local Similarity
RESULT 933
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                                      Best Local Similarity
RESULT 931
                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LIU J.
                                                                                                                        EP1033405-A2.
                                                                                                                                        06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-FEB-2004
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                        Query Match
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RESULT 935

RESULT 939
ID ADX954
DE ADX954
DE Plant
PN US2004
PD 19-FEB
PA (ZHOU/
PA (SCNE/

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Length 102;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 8.2%; Score 96.5; DB 3; Length 621; Local Similarity 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 8.2%; Score 96.5; DB 4; Length 621;
Best Local Similarity 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 195;
                                                                                                                                                                                Score 101; DB 3; Length 214; Pred. No. 0.097;
                                                                                                                                                                                                                                                                                                                                                                      Score 101; DB 3; Length 237; Pred. No. 0.11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8:3%; Score 98; DB 3; Length 208; 21.2%; Pred. No. 0.19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8.3%; Score 98; DB 3; Length 208; 21.2%; Pred. No. 0.19;
                                                                                                                                                                                                                                                               AAG51637 standard; protein; 237 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 65557
EP10313405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAGS4218 standard; protein; 208 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 69107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABB68400 standard; protein; 195 AA.
Drosophila melanogaster polypeptide SEQ ID NO 31992.
WO200171042-A2.
8.6%; Score 101.5; DB 8; 30.7%; Pred. No. 0.03;
                                                              AAGS1638 standard; protein; 214 AA.
Arabidopsis thaliana protein fragment SEQ ID NO:
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAGO4572 standard; protein; 208 AA.
Arabidopsis thaliana protein fragment SEQ ID NO:
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU53138 standard; protein; 544 AA.
Human testes-derived protein from DKFzphtes3_8g5
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8.2%; Score 96.5; DB 4; 21.8%; Pred. No. 0.25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU53359 standard; protein; 537 AA.
Human testes-derived DKFZphtes3_895 homologue.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 8.2%; Score 96.5; Di
Best Local Similarity 24.2%; Pred. No. 1.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB93236 standard; protein; 621 AA.
Human protein sequence SEQ ID NO:12237.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GEHU-) GERMAN HUMAN GENOME PROJECT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GEHU-) GERMAN HUMAN GENOME PROJECT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY83077 standard; protein; 621 AA.
F-box protein FBP-9.
WO200012679-A1.
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Human translocation associated protein designated Gp25L-H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                   DB 5; Length 621;
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                                                                                                                                                                                                                                                            Length 621;
                                                                                                                                                                                                                                                                                                                                                                                                                  .8.2%; Score 96.5; DB 9; Length 621; 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 634;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8.2%; Score 96.5; DB 9; Length 621; 24.2%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 628;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 9; Length 628
                                                                                                                                                                                                                                                                                                               ADX05819 standard; protein; 621 AA.
Cyclin-dependent kinase modulation biomarker SEQ ID NO 384.
WO2005012875-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADXO5821 standard; protein; 628 AA.
Cyclin-dependent kinase modulation biomarker SEQ ID NO 386.
WO2005012875-A2.
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Arabidopsis thaliana protein fragment SEQ ID NO: 13638.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8.2%; Score 96.5; DB 9; 24.2%; Pred. No. 1.4;
                                                                                                                                                                                                                                                          Score 96.5; DB 7;
Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8.2%; Score 96.5; DB 4; 24.2%; Pred. No. 1.4;
                                                                                                                                                       ADB79933 standard; protein; 621 AA.
Human putative F-box protein Fbx21, SEQ ID 173
EP1279744-A2.
                                    PD 18-JUL-2002.
PP (UNYY) UNIV NEW YORK STATE.
Query Match
Best Local Similarity 24.2%; Pred. No. 1.3;
RESULT 950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PA (BRIM') BRISTOL-MYERS SQUIBB CO.
Query Match
Best Local Similarity 24.2%; Pred. No. 1.4;
RESULT 955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8.2%; Score 96.5; Di 24.2%; Pred. No. 1.3;
AAO22463 standard; protein; 621 AA.
Human P-box protein FBP9 SEQ ID No 28.
WO200255665-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AEA23925 standard; protein; 628 AA.
Human PRO polypeptide SEQ ID NO 467.
WO2005051988-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEL92451 standard; protein; 621 AA.
Human F-box protein FBP9.
WO2006113251-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADY62385 standard; protein; 621 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU31647 standard; protein; 634 AA. Novel human secreted protein #2138 WO200179449-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW80293 standard; protein; 218 AA
                                                                                                                             Lative P-box protein;
29-784-A2.
29-78N-2003.
PA (WARN) WARNER LAMBERT CO.
Query Match
Best Local Similarity 24.2%; PrekRSULT 951
ID ADXO5819 standard; profit DE CYCLIN-dependent PN W0200501287F
PD 10-FFP
PA
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(BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYNY ) UNIV NEW YORK STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYNY ) UNIV NEW YORK STATE. (CELG-) CELGENE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human F-box protein, FBP9.
WO2005020922-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-JUN-2005.
(GETH ) GENENTECH INC.
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RESULT 958
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Best Local Similarity
RESULT 954
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-MAR-2005
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Length 107;
                                                                                                                                                                                                                                                                                                                                                                 Length 169;
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                                                                             Length 218;
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Pred. No. 2.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.9%; Score 93; DB 5; Length 493; 20.4%; Pred. No. 2.3;
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Pred. No. 2.3;
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                                                                                                                                                                                                                                                                                    AAG51639 standard; protein; 169 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 65559
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAG49571 standard; protein; 493 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 62725.
EP1033405-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGG28393 standard; protein; 493 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 33594
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAG21783 standard; protein; 493 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 24466.
EP1033405-A2.
06-SEP-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disease treating protein complex-derived protein #374 EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                    vuery Match
Best Local Similarity 21.0%; Pred. No. 0.43;
RESULT 961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 19.8%; Pred. No. 0.62;
RESULT 962
                                                                                                                                                                                                                               7.9%; Score 93.5; DB 5; 23.9%; Pred. No. 0.22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 93.5; DB 7;
Pred. No. 0.62;
PN USS831055-A.
PD 03-NOV-1998.
PA (INCY-) INCYTE PHARM INC.
Query Match
Best Local Similarity 21.4%; Pred. No. 0.54;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Herbicidally active polypeptide SEQ ID NO 1548.
W0200210-02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 93; DB 3;
Pred. No. 2.3;
                                                                                                                                ABP61452 standard; protein; 107 AA.
Human NF-kB activating protein SEQ ID NO 57.
WO200253737-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.8%; Score 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADI16915 standard; protein; 400 AA. Rat NOVX protein homologue SeqID 451 WO200268649-A2.
                                                                                                                                                                                                                                                                                                                                                                                                        Protein sequence #SEQ ID 709.
20-Now Communications of the protein sequence #SEQ ID 709.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADK62496 standard; protein; 219 AA
                                                                                                                                                                                  11-JUL-2002.
(ASAH ) ASAHI KASEI KOGYO KK.
Query Match 7.9%;
Best Local Similarity 23.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.98;
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(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-2003.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-NOV-2002.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 963
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Best Local Similarity
RESULT 965
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RESULT 964
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(FARB ) BAYER AG.
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DB 9; Length 276;
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                                                                                                                                                                                                                 7.6%; Score 89; DB 3; Length 168; 24.3%; Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 89; DB 8; Length 223;
Pred. No. 1.9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4; Length 408;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 5; Length 605;
13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABG93808 standard; protein; 605 AA.
Transcription factor stress-related protein (TFSRP)
US2002102695-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADMZ6073 standard; protein; 1257 AA.
Hyperthermophile Methanopyrus kandleri protein #679.
WO2003076575-A2.
                                                                                                                                                                                                                                                                                 ADY23730 standard; protein; 223 AA.
Plant full length insert polypeptide segid 71514.
US2004034888-A1.
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8
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Pred. No. 5.9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 7.6%; Score 89; DB 4
Local Similarity 37.7%; Pred. No. 4.5;
                                                            7.7%; Score 90.5; DI 24.7%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 87;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 87;
Pred. No.
                                                                                                                  AAB27560 standard; protein; 168 AA. Human secreted protein SEQ ID NO: 61. WQ200055175-A1. 21-SEP-2000. (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human diagnostic protein; 408 AA. WO2001-75067-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW47273 standard; protein; 1946 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAU71799 standard; protein; 605 AA.
Zinc finger-3 (ZF-3).
WO200177311-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADH78770 standard; protein; 384 AA Pea GPA1 ortholog, SEQ ID 50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 7.6%;
Local Similarity 22.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match
Local Similarity 23.0%;
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Best Local Similarity 28.2%;
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PN US2002102695-A1.
PD 01-AUG-2002.
PA (SILV/) SILVA O D C E.
PA (BOHN/) BOHNERT H J.
PA (THIE/) THIELEN N V.
PA (CHEN/) CHEN R.
                                                                                                                                                                                                                                                                                                                                                                (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                             (TABA/) TABASKA J E. (CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                            SCREEN S E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                              Local Similarity
                                                                                                                                                                                                                                          Local Similarity
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(HYSE-) HYSEQ INC.
                    16-DEC-2004.
(RIKE ) RIKEN KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2004003146-A2.
08-JAN-2004.
  WO2004108930-A1.
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                                                            Query Match
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Mouse small endoplasmic reticulum resident protein 3 (smERp3) version #2.
EP1447412-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADV67442 standard; protein; 276 AA.
Amino acid sequence of a protein involved in FGF signal transduction.
                                                                                                                                          DB 6; Length 572;
                                                                                                                                                                                                                                                                                                        DB 7; Length 572;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 572;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 6; Length 218;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 276;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 276;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.7%; Score 91; DB 5; Length 1450; 21.3%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.7%; Score 91; DB 8; Length 372; 24.2%; Pred. No. 2.4;
                                                                                                                                                                                                ADK63196 standard; protein; 572 AA.
Disease treating protein complex-derived protein #724.
EP1338608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA22701 standard, protein, 218 AA.
A. gossypii transport protein Oligo 167 SEQ ID 45.
WO2003014353-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UVUT-) UNIV UTRECHT HOLDING BV.
ry Match
7.7%; Score 90.5; DB 8;
t Local Similarity 24.7%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADH78766 standard; protein; 372 AA.
Nicotiana plumbaginifolia GPA1 ortholog, SEQ ID
WO2004003146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.7%; Score 90.5; Di
24.7%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.7%; Score 90.5; Di 21.4%; Pred. No. 1.3;
  23.0%; Pred. No. 2.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAU02498 standard; protein; 276 AA.
Murine trinucleotide repeat protein (TRP).
WO200130798-A1.
                                                                                                                                                                                                                                                                                                      7.8%; Score 91.5; 22.0%; Pred. No. 4;
                                                                                                                                        7.8%; Score 91.5; 22.0%; Pred. No. 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 7.8%; Score 91.5; Best Local Similarity 22.0%; Pred. No. 4; RESULT 971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (INRG) INRA INST NAT RECH AGRONOMIQUE.
Query Match 7.7%; Score 91;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABB55397 standard; protein; 1450 AA.
Lactococcus lactis protein yvcC.
                                       ABR53173 standard; protein; 572 AA. Protein sequence #SEQ ID 1211.
                                                                                                                                                                                                                                                                                                                                                                ADS43610 standard; protein; 572 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PARA-) PARADIGM GENETICS INC. (UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                 Bacterial polypeptide #22040.
US2003233675-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PA (DELT-) DELTAGEN INC.
Query Match
Best Local Similarity 2
RESULT 975
                                                                                                                                                                                                                                                                                                                                                                                                                          18-DEC-2003.
(CAOY/) CAO Y.
(HINK/) HINKLE G J.
(SLAT/) SLATER S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CHEN/) CHEN X. (GOLD/) GOLDMAN B S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
RESULT 972
                                                                                                                      (CELL-) CELLZOME AG.
Query Match
                                                                                                                                                                                                                                                                                   (CELL-) CELLZOME AG.
Best Local Similarity RESULT 968
                                                                                                                                                         Best Local Similarity RESULT 969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
RESULT 976
                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BADI ) BASF AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FR2807446-A1.
                                                                                EP1258494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JAN-2004.
                                                                                                     20-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-MAY-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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#3

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AEA59839 standard; protein; 690 AA.
Streptococcus pneumoniae ORF amino acid sequence SEQ ID NO:4604.
US2005136404-A1.
                                                                               DB 2; Length 1946;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABU02615 standard; protein; 690 AA.
S. pneumoniae type 4 strain protein from coding region #2193.
03-077021-A2.
03-067-2002.
                                                                                                                                                                                                                                   DB 8; Length 391;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7.3%; Score 86; DB 6; Length 491; 24.5%; Pred. No. 12;
                                                                                                                                                                                                                                                                                                                                                                                    Score 86; DB 4; Length 488;
Pred. No. 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OSCUSTANT OF THERAPEUTICS CORP.

(GENO-) GENOME THERAPEUTICS CORP.

(GENO-) GENOME THERAPEUTICS CORP.

7.3%; Score 86; DB 8; Length 690;

(ery Match

7.3%; Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 86; DB 6; Length 690;
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 86; DB 9; Length 690;
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 86; DB 8; Length 687
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 8; Length 690, 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU22666 standard; protein; 491 AA.
Protein encoded by Prokaryotic essential gene #8193.
WO200277183-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADR95969 standard; protein; 690 AA.
Novel S. pneumoniae protein sequence, SEQ ID 4604.
US6800744-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM92253 standard; protein; 687 AA.
S pneumoniae antigenic protein sequence SeqID450.
WO20040010609-A2.
(TUFT ) UNIV TUFTS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Streptococcus pneumoniae protein, Seq ID No 4221.
US6699703-B1.
                                                                                                                                                                                                                               Query Match 7.3%; Score 86; DB 8 Best Local Similarity 22.1%; Pred. No. 8.8; RESULT 986
                   PD 03-DEC-1997.
PA (NEST ) SOC PROD NESTLE SA.
Query Match
Best Local Similarity 22.5%; Pred. No. 79;
RESULT 985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-0CT-2004.
(GENO-) GENOME THERAPEUTICS CORP.
(GENO-) GANOME THERAPEUTICS CORP.
(ery Match
7.3%; Score 86;
Lactobacillus bulgaricus protease P.
                                                                                                                                                                                                                                                                       Novel bone marrow polypeptide #46.09-ANTS-200157187-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADK47706 standard; protein; 690 AA.
                                                                                                                            AD065925 standard; protein; 391 AA.
Novel human protein sequence #898.
EP1440981-A2.
                                                                                                                                                                                          28-JUL-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                  7.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DOUC/) DOUCETTE-STAMM L A. (BUSH/) BUSH D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHIR-) CHIRON SPA.
(GENO-) INST GENOMIC RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-OCT-2002.
(ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
RESULT 987
ID ABU2266 standard; px
DE Protein encoded by Px
PN W020277183-A2.
PD 03-OCT-2002.
PA (ELIT-) ELITRA PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 988
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Best Local Similarity
RESULT 991
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Best Local Similarity
RESULT 992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
RESULT 989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 990
                                                                                                                                                                                                                                                                                                                                             09-AUG-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JUN-2005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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Length 690;
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Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 86; DB 10; Length 690;
Pred. No. 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.3%; Score 86; DB 6; Length 2301;
19.3%; Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 86; DB 2; Length 2938;
Pred. No. 1.6e+02;
                                                                                                                                                                                                                                           Length 690;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 690
AEJ71634 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7074914-B1.
                                                                                                                                         AEJ78609 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7081530-B1.
                                                                                                                                                                                                                                                                                              AEJ85969 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7098023-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                AEL08288 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7115731-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AEL15538 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221.
US7122368-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AEL53946 standard; protein; 690 AA.
Streptococcus pneumoniae strain 14453 protein, SEQ:4221
                                                                                             DB 10;
20;
                                                                                                                                                                                                                                             DB 10;
20;
                                                                                                                                                                                                                                                                                                                                                                                            ; DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 10;
20;
                                           PD 11-JUL-2006.
PA (SNFI) SANOFI PASTEUR LIEE.
Query Match
7.3%; Score 86;
Rest Local Similarity 22.3%; Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                              Score 86;
Pred. No.
                                                                                                                                                                                                                                               Score 86;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 86;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABUS2379 standard; protein; 2301 AA.
Human GPCR related protein NOV29a.
WO200279398-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR59923 standard; protein; 2938 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADN18817 standard; protein; 3092 AA
                                                                                                                                                                                               PD 25-JUL-2006.
PA (SNFI ) SANOFI PASTEUR LIEE.
QUERY, MATCh
Best Local Similarity 22.3%;
RESULT 995
                                                                                                                                                                                                                                                                                                                                                  29-AUG-2006.
(SNFI ) SANOFI PASTEUR LTD.
7.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 22.3%;
RESULT 998
                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 22.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (SNPI ) SANOFI PASTEUR LTD.

17.3%;

1 Local Similarity 22.3%;
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Local Similarity 22.3%;
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Best Local Similarity 19.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bacterial polypeptide #1470.
US2003233675-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33.78;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-OCT-2006.
(SNFI ) SANOFI PASTEUR LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-OCT-2006.
(SNFI ) SANOFI PASTEUR LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SCHE ) SCHERING CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-OCT-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CAO Y.
HINKLE G J.
SLATER S C.
CHEN X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SLAT/) SLATER S C.
(CHEN/) CHEN X.
(GOLD/) GOLDMAN B S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAP protein Iral.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 997
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22-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABUS3080 standard; protein; 711 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #13.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                AABS6790 standard; protein; 173 AA.
Human prostate cancer antigen protein sequence SEQ ID NO:1368.
WO200055174-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 384;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.3%; Score 85.5; DB 3; Length 384; 19.4%; Pred. No. 9.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.3%; Score 85.5; DB 8; Length 601; 23.0%; Pred. No. 18;
                                                                                                                                                                                                                                                                                  DB 3; Length 147;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 384;
                                                                                                                                   7.3%; Score 85.5; DB 3; Length 147; 25.2%; Pred. No. 2.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 5; Length 623;
   Length 3092;
                                                                                                                                                                                         AAB27611 standard; protein; 147 AA.
Human secreted protein BLAST search protein SEQ ID NO: 112.
WO200055175-A1.
                                                    AAB27610 standard; protein; 147 AA.
Human secreted protein BLAST search protein SEQ ID NO: 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAG21784 standard; protein; 384 AA..
Arabidopsis thaliana protein fragment SEQ ID NO: 24467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAG49572 standard; protein; 384 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 62726
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAG28394 standard; protein; 384 AA.
Arabidopsis thaliana protein fragment SEQ ID NO: 33595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 7.3%; Score 85.5; DB 3;
Best Local Similarity 19.4%; Pred. No. 9.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                       7.3%; Score 85.5; DB 3; 25.2%; Pred. No. 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.3%; Score 85.5; DB 3; 19.4%; Pred. No. 9.6;
 7.3%; Score 86; DB 8; I
33.7%; Pred. No. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADW07163 standard; protein; 601 AA.
Bos taurus keratin 5 protein sequence SeqID3
NZS25023-A.
                                                                                                                                                                                                                                      G1-SEP-2000.

(HUMA-) HUMAN GENOME SCI INC.

7.3%; Score 85.5; Di
ery Match
7.3%; Pred. No. 2.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.3%; Score 85.5; I
18.2%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABB77434 standard; protein; 623 AA. Human tumour marker protein se20-7.WO200238803-A2.
                                                                                  WO200055175-A1.
21-SEP-2000.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (STAN/) STHERLAND G T.
(STAN/) STANFIELD A M.
(SPEL/) SPELMAN R J.
(ANKEA) ANKERSMIT-UDY A E L.
                                                                                                                                                                                                                                                                                                                                                                                                     (HUMA+) HUMAN GENOME SCI INC. (ROSE/) ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Luery Match
Best Local Similarity 1.
RESULT 1010
ID ABU53080 Str.
DE INTRAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
            Best Local Similarity RESULT 1002
                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNEL/) SNELL R G. FORD/) FORD C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EP1033405-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match
                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
   Query Match
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                                                                                                                                                                           RESULT 1003
                                                                                                                                                                                                                                                                                                                  RESULT 1004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BEEE
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ABUS3072 standard; protein; 941 AA.
Intracellular trafficking-associated DKFZphtes3_1913 homologue #5.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                      ABU53070 standard; protein; 951 AA.
Intracellular trafficking-associated DKF2phtes3_1913 homologue #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABUDSJO71 standard; protein; 953 AA.
Intracellular trafficking-associated DKFZphtes3_1913 homologue #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            авибзибе standard; protein; 953 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #2.
WO200112659-A2.
                                                                        ABUSS073 standard; protein; 907 AA.
Intracellular trafficking-associated DKFZphtes3_1g13 homologue #6.
WO200112659-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 2230;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2228;
                                                                                                                                                                      Score 85.5; DB 4; Length 907; Pred. No. 33;
                Length 711;
                                                                                                                                                                                                                                                                                                                                   Length 941;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 951;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU07445 standard; protein; 2230 AA.
Protein differentially regulated in prostate cancer #48
                    DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 7.3%; Score 85.5; DB 6; Best Local Similarity 18.2%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.3%; Score 85.5; DB 7;
18.2%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                   DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 85.5; DB 7;
Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR61600 standard; protein; 2230 AA.
Human golgin-245 splice variant 2 polypeptide.
WO2003087403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR61601 standard; protein; 2250 AA.
Human golgin-245 splice variant 3 polypeptide.
WO2003087403-A2.
ACCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR61599 standard; protein; 2228 AA.
Human golgin-245 splice variant 1 polypeptide
WO2003087403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB
                Score 85.5; I
Pred. No. 23;
                                                                                                                                                                                                                                                                                                                           Best Local Similarity 18.2%; Pred. No. 35; RESULT 1013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 85.5; Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 85.5;
Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 85.5;
Pred. No. 36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GEHU-) GERMAN HUMAN GENOME PROJECT.
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                                                                                                                               (GEHU-) GERMAN HUMAN GENOME PROJECT.
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GERMAN HUMAN GENOME PROJECT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (EVOT-) EVOTEC NEUROSCIENCES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ORIG-) ORIGENE TECHNOLOGIES INC.
                  Best Local Similarity 18.2%;
RESULT 1011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 7.3%;
Best Local Similarity 18.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 7.3%;
Best Local Similarity 18.2%;
RESULT 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 18.2%;
RESULT 1017
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 7.3%;
Best Local Similarity 18.2%;
                                                                                                                                                                            Query Match 7.3%;
Best Local Similarity 18.2%;
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                                                                                                                                                                                                                                                                                               22-FEB-2001
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DB 8; Length 773;

Length 227;

9

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Score 84; DB 4; Length 1605;
Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Match 7.1%; Score 84; DB 5; Length 486; Local Similarity 20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       'Match 7.1%; Score 84; DB 8; Length 486; Local Similarity 20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADV88989 standard; protein; 486 AA.
Streptococcus agalactiae protein sequence, SEQ ID 1383
                                                                                                                                                 Alloicoccus otitis antigenic protein SEQ ID NO:6554 WO2001048304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila melanogaster polypeptide SEQ ID NO 37917.
WO200171042-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC31334 standard; protein; 827 AA.
Human novel polypeptide sequence, SEQ ID NO:1416.
WO2003029271-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          y Match 7.1%; Score 84; DB 8;
Local Similarity 20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.1%; Score 84; DB 8;
20.5%; Pred. No. 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADV80242 standard; protein; 486 AA.
Streptococcus agalactiae protein, SEQ ID 1383
WO200292818-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Streptococcus agalactiae protein, 486 AA.
WO2002918-A2.
                                                                                                                                                                                                                                                         Query Match 7.1%; Score 84; DB (Best Local Similarity 19.6%; Pred. No. 6.5; RESULT 1029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7.1%; Score 84; DB 32.2%; Pred. No. 42;
                                                                                                        Score 84.5; 1
Pred. No. 34;
                                                                                                                                                                                                                                                                                                                         ABP27811 standard; protein; 486 AA.
Streptococcus polypeptide SEQ ID NO 4798.
WO200234771-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADL24093 standard; protein; 2301 AA.
Human NOVX polypeptide #69.
US2004002120-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 1605 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-NOV-2002.
(INSP ) INST PASTEUR.
(CNRS ) CNRS CENT NAT ECH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (INSP ) INST PASTEUR.
(CNRS ) CNRS CENT NAT RECH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-NOV-2002.
(INSP ) INST PASTEUR.
(CNRS ) CNRS CENT NAT RECH SCI.
                                                                                                        7.2%;
                                                                                                                                                                                                                           12-JUN-2003.
(AMHP ) WYETH HOLDINGS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.1%;
                                                                                                                                                                                                                                                                                                                                                                                  02-MAY-2002.
(CHIR-) CHIRON SPA.
(GENO-) INST GENOMIC RES.
                                                               (CHEN/) CHEN X. (GOLD/) GOLDMAN B S.
                        HINKLE G J. SLATER S C.
                                                                                                  Query Match
Best Local Similarity
RESULT 1028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1033
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-SEP-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABB70375 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JAN-2004.
(KEKU/) KEKUDA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FR2824074-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                 (SLAT/
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 7.3%; Score 85.5; DB 8; Length 4447;
Best Local Similarity 24.4%; Pred. No. 3.3e+02;
RESULT 1022
                        Length 2250;
                                                                                                                                                                                  Length 2252;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.2%; Score 84.5; DB 7; Length 773; 21.1%; Pred. No. 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.2%; Score 85; DB 8; Length 223; 22.0%; Pred. No. 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 419;
12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADK63862 standard; protein; 773 AA.
Disease treating protein complex-derived protein #1444.
EP1338608-A2.
                                                                                                                    PD 23-OCT-2003.
PA (EVOT-) EVOTEC NEUROSCIENCES GMBH.
Query Match 7.3%; Score 85.5; DB 7;
Best Local Similarity 18.2%; Pred. No. 1.2e+02;
RESULT 1021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADY10508 standard; protein; 223 AA.
Plant full length insert polypeptide seqid 66323.
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       wuery Match 7.2%; Score 84.5; DB 6;
Best Local Similarity 21.1%; Pred. No. 34;
RESULT 1026
Query Match 7.3%; Score 85.5; DB 7; Best Local Similarity 18.2%; Pred. No. 1.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PD 03-APR-2003.

PA (INCY-) INCYTE GENOMICS INC.

Query Match 7.2%; Score 85; DB 6;

Best Local Similarity 32.2%; Pred. No. 27;
                                                                           ABR61602 standard; protein; 2252 AA.
Human golgin-245 splice variant 4 polypeptide.
WO2003087403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.2%; Score 85; 22.7%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABG13105 standard; protein; 419 AA.
Novel human diagnostic protein #13096.
WO200175067-A2.
                                                                                                                                                                                                                                         ADS44713 standard, protein; 4447 AA.
Bacterial polypeptide #23143.
US2003233675-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR69620 standard; protein; 722 AA.
Human CGDD-20 protein.
WO2003027263-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR53767 standard; protein; 773 AA.
Protein sequence #SEQ ID 2399.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADN18818 standard; protein; 773 AA.
Bacterial polypeptide #1471.
US2003233675-A1.
18-DEC-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lu-) CELLZOME AG.
Lu-Y Match
Best Local Similarity 2
RESULT 1027
ID ADN18918 stand-
DE Bacterial
PN US2n-
PD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-NOV-2002.
(CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                (CHEN/) CHEN X.
(GOLD/) GOLDMAN B S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 1024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 1023
                                                                                                                                                                                                                                                                                                                       (CAOY/) CAO Y.
(HINK/) HINKLE G J.
(SLAT/) SLATER S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-OCT-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP1258494-A1.
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Length 486;

Length 486;

Length 827;

7;

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Human protein encoded by clone THYMU20098350.
EP1308459-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UNIV NORTH CAROLINA STATE.

7.1%; Score 83.5; DB 9; Length 406; Similarity 18.2%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.1%; Score 83.5; DB 4; Length 399; 19.7%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 8; Length 326;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 8; Length 2301;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB75224 standard; protein; 399 AA.
Drosophila gustatory receptor GR59E.2 protein sequence.
WO200077208-A2.
21-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADX96054 standard, protein; 326 AA.
Plant full length insert polypeptide segid 58718
US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 7.1%; Score 84; DB 8; 1
Best Local Similarity 18.3%; Pred. No. 1.8e+02;
RESULT 1036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAE30522 standard; protein; 399 AA.
Fruit fly gustatory receptor protein, Gr59E2.
W0200268593-A2.
06-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.1%; Score 83.5; I 21.3%; Pred. No. 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7.1%; Score 83.5; 19.7%; Pred. No. 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEC57546 standard; protein; 406 AA.
L. acidophilus ABC transporter permease.,
WO2005084411-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB65618 standard; protein; 567 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYCO ) UNIV COLUMBIA NEW YORK.
                                                                                                                                       GORMAN L.
MALYANKAR U M.
BOLDOG F L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SHIMKETS R A.
RASTELLI L.
SPADERNA S K.
LAROCHELLE W J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KHRAMTSOV N V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          FERNANDES E R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HERRMANN J L.
      TCHERNEV V T.
                                                                                                     FRNET C A M
                                                                                                                                                                                                                    SHENOY S G.
PADIGARU M.
TAUPIER R J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
RESULT 1038
                                                                                                                                                                                                                                                                                MILLER C E. CASMAN S J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 1039
                                                                                                                                                                                                                                                                                                                     PENA C E A
                                                                                                                                                                                                                                                                                                                                                                                 SMITHSON G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYYA ) UNIV YALE.
                                                                                                                                                                                                                                                                                                                                                             GUSEV V Y.
                                                                                                                                                                                                                        SHEN/
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AEBS6515 standard; protein; 1022 AA.
Radiochemotherapy response detection associated protein SEQ ID NO 90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 9; Length 1022;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 8; Length 590;
                                                                                              Length 567;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 7.1%; Score 83.5; DB 4; Length 914;
Best Local Similarity 19.7%; Pred. No. 55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 8; Length 590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ37107 standard; protein; 590 AA.

Human malignant pleural mesothelioma (MPM) protein #6.
US2003129760-A1.

(BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila melanogaeter polypeptide SEQ ID NO 31677. W0200171042-A2. 27-SEP-2001. (PREE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 9;
                                                                                              DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 7.1%; Score 83.5; DB 9;
Best Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (IPSO-) IPSOGEN.
(INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
(PAOL-) INST PAOLI CALMETTES.
                                                                                          Query Match 7.1%; Score 83.5; I
Best Local Similarity 22.5%; Pred. No. 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MILL-) MILLENNIUM PHARM INC.
Query Match 7.1%; Score 83.5; I
Best Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 7.1%; Score 83.5; I
Best Local Similarity 25.1%; Pred. No. 64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 7.1%; Score 83.5; 1
Best Local Similarity 22.5%; Pred. No. 29;
RESULT 1042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 7.1%; Score 83.5; Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 7.1%; Score 83.5;
Best Local Similarity 22.5%; Pred. No. 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 22.5%; Pred. No 22.5%; P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AEB87738 standard; protein; 590 AA.
Human cytokeratin 5, breast tumor marker.
WO2005071419-A2.
                                                                                                                                                                                                                                                                                                                                                                   13-MAR-2003.
(BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADR99136 standard; protein; 590 AA.
Keratin 5, KRT5, SEQ ID 142.
WO2004078035-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB75394 standard; protein; 590 AA.
Prostate cancer marker protein.
WO2003009814-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEA15495 standard, protein; 590 AA.
Human polypeptide #114.
WO2005047534-A2.
                                                                                                                                                                                                                              protein; 590 AA
(HELI-) HELIX RES INST. (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (FARB ) BAYER HEALTHCARE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (FARB ) BAYER PHARM CORP
                                                                                                                                                                                                                              ADB70328 standard; prot.
Keratin 5 SEQ ID NO:20.
WO2003021229-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 06-FEB-2003
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14-APR-2005
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                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
RESULT 1057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADR49442 standard; protein; 276 AA.
Mouse small endoplasmic reticulum resident protein 3 (smERp3) version #1.
EP1447412-A1.
                                                                                                                                                                     AELS7159 standard; protein; 1062 AA.
Human SLIT-ROBO Rho GTPase activating protein 1, SEQ ID NO: 1622.
US2006216722-A1.
28-SEP-2006.
          ADA13347 standard; protein; 1062 AA.
Human intracellular signalling molecule INTSIG-29, SEQ ID NO:29.
WO2003031558-A2.
                                                                                                                                                                                                                                                                                                                                                      7.1%; Score 83.5; DB 10; Length 1062; 25.1%; Pred. No. 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AED11485 standard; protein; 181 AA.
Human gene 16 encoded secreted protein HIGAN47, SEQ ID NO: 46.
US2005214786-A1.
                                                                                                                  DB 6; Length 1062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADJ92059 standard; protein; 474 AA.
Soybean hydroperoxide lyase from clone sdp3c.pk017.j17:fis.
US2004010822-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 7; Length 181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.1%; Score 83; DB 8; Length 282; 24.4%; Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 83; DB 8; Length 474; Pred. No. 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7.1%; Score 83; DB 8; Length 276; 25.0%; Pred. No. 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 83; DB 8; Length 429; Pred. No. 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADY23081 standard; protein; 282 AA.
Plant full length insert polypeptide segid 70865.
US2004034888-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 9; 1
                                                                                                                7.1%; Score 83.5; 1
25.1%; Pred. No. 68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-SEP-2003.
(HUMA-) HUMAN GENOME SCI INC.
(HT Match 7.1%; Score 83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7.1%; Score 83; 23.4%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                             ADC78239 standard; protein; 181 AA.
Human secreted protein SEQ ID NO:46.
WO2003072761-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADTS8337 standard; protein; 429 AA Plant polypeptide, SEQ ID 8414. US2004216190-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-AUG-2004. (UYUT-) UNIV UTRECHT HOLDING BV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-SEP-2005.
(HUMA-) HUMAN GENOME SCI INC.
                                                                       17-APR-2003.
(INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.1%;
                                                                                                                                                                                                                                                  (BETS/) BETSHOLTZ C.
(TRYG/) TRYGGVASON K.
(TAKE/) TAKEMOTO M.
(HELL/) HE L.
(PATR/) PATRAKKAS J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
(SCRE) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
SULT 1052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JAN-2004.
(MCGO/) MCGONIGLE B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
RESULT 1053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
RESULT 1055
                                                                                                                              Best Local Similarity
RESULT 1050
                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 1051
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Best Local Similarity
RESULT 1054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 1056
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
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DB 8; Length 325;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 9; Length 403;
                                                                                                   Length 474;
                                                                                                                                                                                                                                                                                                                                                                      Score 83; DB 8; Length 489;
Pred. No. 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              7.0%; Score 82; DB 6; Length 211; 23.6%; Pred. No. 9.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.0%; Score 82; DB 7; Length 211; 23.6%; Pred. No. 9.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse squalene epoxidase (SPF) protein sequence SeqID2 JP2005095162-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK62500 standard; protein; 211 AA.
Disease treating protein complex-derived protein #376.
                                                                                                                    Best Local Similarity 24.5%; Fred. No. 24;
RESULT 1058

ID ADX78603 standard; protein; 489 AA.

BE Plant full length insert polypeptide seqid 47969.

PN US2004034888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB70282 standard; protein; 984 AA.
C. neoformans amino acid sequence SEQ ID NO:3326.
WO2003052076-A2.
                                                                                                      9;
                                                                                                   ; DB
                                                                                                                                                                                                                                                                                                                                                                                                                            ADP04221 standard; protein; 325 AA.
Human colon specific protein SEQ ID NO:202.
WO2004050858-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Luciy Macch
Beet Local Similarity 20.5%; Pred. No. 77;
RESULT 1063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 82.5; Pred. No. 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 82.5; I
Pred. No. 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 82.5; Pred. No. 21;
AED00636 standard; protein; 474 AA.
Soybean hydroperoxide lyase SEQ ID NO:11.
29-SEP-2005.
(DUPO ) DU PONT DE NEMOURS & CO E I.
ery Match
7.1%; Score 83; D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Murine NOVX protein homologue SeqID 450.
WO200268649-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI16914 standard; protein; 403 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABR5924 standard; protein; 211 AA.
Protein sequence #SEQ ID 713.
EP1258494-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 403 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADH78758 standard; protein; 366 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7.0%;
                                                                                                                                                                                                                                                                                                                                                                        7.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 7.0%;
Best Local Similarity 22.6%;
RESULT 1060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 7.0%;
Best Local Similarity 22.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ARAI/) ARAI H.
(SHIB/) SHIBATA N.
(CHUS ) CHUGAI PHARM CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ELIT-) ELITRA PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-JUN-2004.
(DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-SEP-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                             (LIUJ/) LIU J.
(ZHOU/) ZHOU Y.
(KOVA/) KOVALIC D K.
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                     (SCRE/) SCREEN S E.
(TABA/) TABASKA J E.
(CAOY/) CAO Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CELL-) CELLZOME AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 1065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADZ14402 standard;
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Human PRO872 polypeptide. US2002103125-A1.

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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                            DB 6; Length 610; 44;
                                                                                                                                                                                                                                                                              DB 6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                    Length 610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 6; Length 610;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 6; Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 610;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      ABU96174 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003036144-A1.
                                                                                               Novel human secreted and transmembrane protein PRO872. US2003036147-Al.
                                                                                                                                                                                                                   ABR68121 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027264-A1.
06-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR74888 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040056-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABR94650 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003044926-A1.
                                                                                                                                                                                                                                                                                                                                                                                                 DB 6;
44;
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Human secreted/transmembrane protein (PRO) #46.
US2003044923-Al.
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Human secreted/transmembrane protein (PRO) #46.
22003040062-A1.
27-FEB-2003.
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44;
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Human secreted/transmembrane protein, #41.
US2002160384-A1.
                                                                                                                                                                            Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
      Human PRO polypeptide #46.
US2003032113-Al.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1088
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RESULT 1094
ID ABU13882 standard; r
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RESULT 1087
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RESULT 1086
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RESULT 1089
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RESULT 1091
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RESULT 1084
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Score 82; DB 6; Length 610;
Pred. No. 44;
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Pred. No. 44;
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                                                                                                                                                                                     Novel human secreted and transmembrane protein PRO872. US2003013153-A1.
                                                                                                                                                                                                                                                          ABUD/1998 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.US2003017544-A1.
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Novel human secreted and transmembrane protein PRO872.
US2003003531-A1.
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Human secreted/transmembrane protein (PRO) #46.
US200101616162-A1.
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44;
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Pred. No. 4
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Pred. No.
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Pred. No.
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US2003036141-Al.
20-FEB-2003.
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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                                                              ABU85623 standard; proteir
Human PRO polypeptide #46.
US2003036140-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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            (GETH ) GENENTECH LTD.
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US2003036137-A1.
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Best Local Similarity
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RESULT 1098
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Best Local Similarity
RESULT 1097
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01-AUG-2002
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(GETH ) GENENTECH INC. Best Local Similarity RESULT 1117 Best Local Similarity RESULT 1118 Best Local Similarity Best Local Similarity RESULT 1115 Best Local Similarity RESULT 1116 ABO06194 standard; Best Local Similarity Best Local Similarity Best Local Similarity Local Similarity 27-FEB-2003. 20-FEB-2003. 20-FEB-2003. 30-JAN-2003. 06-FEB-2003. Ouery Match Query Match Query Match Query Match Query Match Query Match Query Match Ouery Match Query Match 6; Length 610; 6; Length 610; DB 6; Length 610; 44; Score 82; DB 6; Length 610; Pred. No. 44; DB 6; Length 610; 44; Length 610; Length 610; 6; Length 610; Length 610; Length 610, ABO00147 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003032101-A1. ABU85008 standard; protein; 610 AA. Novel human secreted and transmembrane protein PRO872. US2003032114-A1. ABR98787 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040064-A1. 27-FEB-2003. ABR78272 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003054474-A1. ABR92210 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003036160-A1. ABR99397 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92 LT 1113
AB002124 standard, protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003040054-A1.
27-FEB-2003. AB016310 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003027267-A1. ABO18851 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003044925-A1. ABO11479 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003036124-A1. 20-FEB-2003. 9 .. 9 . 9 DB 44; 26.3%; Pred. No. 44; Score 82; Pred. No. 4 Score 82; Pred. No. Score 82; Pred. No. Score 82; Pred: No. Score 82; Pred. No. 7.0%; 7.0%; 7.0%; 26.3%; 7.0%; 7.0%; 7.0%; 7.0%; 7.0%; 7.0%; 7.0%; 20-MAR-2003. (GETH ) GENENTECH INC. Query Match Best Local Similarity RESULT 1110 Query Match Best Local Similarity Best Local Similarity RESULT 1104 Query Match Best Local Similarity Best Local Similarity RESULT 1108 Best Local Similarity Best Local Similarity Best Local Similarity RESULT 1114 Best Local Similarity Best Local Similarity Best Local Similarity US2003040063-A1. 27-FEB-2003. 06-FEB-2003. 13-FEB-2003. 20-FEB-2003. 06-MAR-2003. 13-FEB-2003. Query Match Query Match Query Match Query Match Query Match Query Match Query Match

RESULT 1109

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RESULT 1107

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DB 6; Length 610; 44;
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ncoccoys standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003036133-A1.
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US2003022294-A1.
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Novel human secreted and transmembrane protein PRO872
US2003036118-A1.
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Human secreted/transmembrane protein, SEQ ID 92.
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US2003036123-A1.
20-PEB-2003.
                                                                                                       Human secreted/transmembrane protein (PRO) #46.
US2003036134-A1.
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44;
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Human secreted/transmembrane protein (PRO) #46.
US2003040060-A1.
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Human secreted/transmembrane protein (PRO) #46
US2003044916-A1.
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Query Match

RESULT 1126

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US2003008353-A1.
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(GETH ) GENENTECH INC.
    (GETH ) GENENTECH INC.
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RESULT 1140
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               Query Match
Best Local Similarity
RESULT 1135
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Best Local Similarity
RESULT 1138
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RESULT 1142
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                            Length 610;
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Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABU95254 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003036117-A1.
                                                                                                                                                                                                                                                                               ABR67097 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027266-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR70008 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032138-A1.
13-FEB-2003.
                              Score 82; DB 6;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                    ABO15700 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003054483-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6;
44;
                                                                                                                                 Score 82; DB 6;
Pred. No. 44;
                                                                                                                                                                    Human secreted/transmembrane protein (PRO) #46 US2003036128-A1.
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Human secreted/transmembrane protein, PRO872
US2003022298-A1.
                                                                                                                                                                                                                                      7.0%; Score 82; 26.3%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           7.0%; Score 82; 26.3%; Pred. No.
                                                               Human PRO polypeptide #46.
US2003032117-A1.
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US2003032130-Al.
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Human PRO polypeptide #46.
US2003032102-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABU71157 standard; protein; 610 AA
                            7.0%;
                                                                                                                                 7.0%;
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(GETH ) GENENTECH INC.
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US2003036143-A1.
                                                                                                                               Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1127
                                       Best Local Similarity
RESULT 1125
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Best Local Similarity
RESULT 1130
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Best Local Similarity
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RESULT 1133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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RESULT 1132
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US2003036156-A1
20-FEB-2003.
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Query Match

RESULT 1129

D M E C

RESULT 1128

Query Match

RESULT 1131

Query Match

Query Match

Query Match

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Score 82; DB 6; Length 610;
Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
                                                                                                                          DB 6; Length 610;
44;
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44;
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44;
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44;
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                                               ABR69341 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036132-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR60081 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032137-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR67816 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027259-A1.
06-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR65204 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027268-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR68426 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027274-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR71838 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032135-A1.
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44;
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44;
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Human secreted/transmembrane protein, #41.
US2003027162-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7.0%; Score 82; 26.3%; Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                            Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                        ABU81284 standard; protein; 610 AA.
Human PRO polypeptide #46.
US2003017542-A1.
                                                                                                                                                                                 protein; 610 AA
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Pred. No. 44;

26.3%;

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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                     7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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26.3%; Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                     Novel human secreted and transmembrane protein PRO872. US2003032108-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABU93654 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
                                                                                                                                                                                                                                                                                         ABU94944 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
13-FBB-2003:
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US20030-2563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 26.3%; Pred. No. 44; RESULT 1153
ID ABR68731 standard; protein; 610 AA.
DE Human secreted polypeptide PRO872, SEQ ID NO:92.PN US2003027271-A1.
                                                                                                                                                                                                                                                                                                                                                            DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 26.3%; Score 82; DB 6; RESULT 1149
ID ABUG4003 standard; protein; 610 AA.
DE Human secreted/transmembrane protein (PRO) #46.
PD 13-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABU83088 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032105-A1.
                                                 ABU89008 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003022297-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                              Score 82;
Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO25913 standard; protein; 610 AA.
Human PRO872 polypeptide.
US2002127576-A1.
12-SEP-2002.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                              7.0%;
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RESULT 1151
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26.3%;
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13-FEB-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 1150
ID ABU93654 standard; pr
DE Novel human secreted
PP US2003032119-A1.
PD 13-FEB-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 1152
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1147
                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 1148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 1154
                                                                                                                                    Best Local Similarity
RESULT 1146
                                                                                                                                                                                                                          13-FEB-2003.
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                                                                                                    30-JAN-2003
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                                                                                                                     Query Match
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                                   RESULT 1145
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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                            Length 610;
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                                                                                                                                                                                                        ABU85928 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003022300-Ai.
                                                                                                                                                                                                                                                                                                 ABU82215 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003036136-A1.
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Novel human secreted and transmembrane protein PRO872.
US2003032104-A1.
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                         ABR99092 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
USO03040068-A1.
27-FEB-2003.
                                                                            DB 6;
44;
                                                                                                                                                                     DB 6;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted/transmembrane protein (PRO) #46.
US2003032109-A1.
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44;
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                              Score 82;
Pred. No.
                                                                                                          Human PRO polypeptide #46. US2003027280-Al.
                                                                                                                                                                                                                                                                                                                                                                                               protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABU83698 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU93964 standard; protein; 610 AA
                                                                            7.0%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Human PRO polypeptide #46.
US203036138-Al.
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Human PRO polypeptide #46.
US2003040066-Al.
Best Local Similarity RESULT 1155
                                                                                         Best Local Similarity RESULT 1156
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                           20-FEB-2003.
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                                                                                                                                                                       Query Match
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                                                                              Query Match
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Score 82; DB 6; Length 610;
Pred. No. 44;
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                                              Length 610;
                                                                                                                                                                                                                                                                                                                                                                Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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44;
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Pred. No. 44;
                                                                                                  Novel human secreted and transmembrane protein PRO872. US2003022296-A1.
Novel human secreted and transmembrane protein PRO872
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADUSDUS Standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US.20132131-A1.
                                                                                                                                                                                                                                                                                                    ABR90905 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040058-A1.
                                                                                                                                                                                     ABR66487 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027281-A1.
                                              .
9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted/transmembrane protein; 610 AA.
US2003032129-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                 DB 6;
                                     DB 6
                                                                                                                                                                                                                                                                                                                                                                                                       ABUS8922 standard; protein; 610 AA.
Human sectreted/transmembrane protein, #41.
US2002142921-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                              Score 82; DB
Pred. No. 44;
                                           Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                7.0%; Score 82; 26.3%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 82;
Pred. No.
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US2003017540-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human PRO polypeptide #46.
US2003032106-A1.
                                                                                    protein; 610 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU94637 standard, protein, 610 AA.
Human PRO polypeptide #46.
US2003032103-A1.
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Human PRO polypeptide #46.
                                                                                                                                             7.0%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
                                                                                                                                                         Best Local Similarity
RESULT 1167
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Best Local Similarity
RESULT 1168
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Best Local Similarity
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Best Local Similarity
RESULT 1171
                                                     Best Local Similarity
RESULT 1166
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Best Local Similarity
RESULT 1172
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Best Local Similarity
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Best Local Similarity
                                                                                   ABU99817 standard;
              US2003036155-A1.
                            20-FEB-2003.
                                                                                                                               30-JAN-2003.
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                                                                                                                                             Query Match
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ID ABI
DE NO
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Length 610;
                          Length 610;
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44;
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44;
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44;
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                                                                                                                                           ABU92300 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003022187-A1.
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Novel human secreted and transmembrane protein PRO872.
US2003036154-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU59365 standard; protein; 610 AA.
Novel human secreted or transmembrane protein PRO1063.
US2003027985-A1.
                                           Human secreted polypeptide PRO872, SEQ ID NO:92. US2003032139-Al. 13-PEB-2003.
                                                                                                                                                                                                                                                                                                                               ABR65877 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003036165-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR64594 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003027262-A1.
                        DB 6;
44;
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44;
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Human secreted/transmembrane protein (PRO) #46
US2003036142-A1.
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44;
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44;
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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US2003032110-A1.
                                                                                                                                                                                                                                           protein; 610 AA
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                        7.0%;
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US2003022301-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                    Best Local Similarity
                                                                                                                                                                                                                                           ABU98478 standard;
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RESULT 1182
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US2003032107-A1.
13-FEB-2003.
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                        Query Match
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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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44;
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44;
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44;
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Pred. No. 44;
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44;
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Pred. No. 44;
                                                                                                           7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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Novel human secreted and transmembrane protein PRO872.
23-JAN-2003.
                                    ABU90182 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
US2003036153-A1.
20-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR69703 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032122-A1.
                                                                                                                                                                                                                                                                                                                                                                         ABR70923 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003040069-A1.
                                                                                                                                                                                                                                               RESULT 1188
ID ABO10869 standard; protein; 610 AA.
DB Human secreted/transmembrane protein (PRO) #46.
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Human secreted/transmembrane protein (PRO) #46.
US2003032116-A1.
                                                                                                                                                    AB003597 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044931-A1.
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Pred. No. 4
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Pred. No. 4
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Pred. No. 4
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Pred. No.
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Pred. No.
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US2003032128-Al.
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Human PRO protein #46.
US2003036139-A1.
20-FEB-2003.
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Best Local Similarity
RESULT 1193
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RESULT 1195
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Best Local Similarity
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Best Local Similarity
RESULT 1192
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RESULT 1196
               Best Local Similarity
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                               RESULT 1186
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Length 610;
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Novel human secreted and transmembrane protein PRO872.
US2002177164-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABU96773 standard; protein; 610 AA.
Novel human secreted and transmembrane protein PRO872.
13-2003032140-A1.
113-PEB-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR70618 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040076-A1.
                                                                                                                                                                                   Human secreted/transmembrane protein (PRO) #46.
US20031652-A1.
                                                        DB 6;
44;
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44;
                                                                                                            Human secreted/transmembrane protein (PRO) #46. US2003017543-A1.
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44;
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Human secreted/transmembrane protein #46.
US2002127584-A1.
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Pred. No.
                                                        Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
ABU93349 standard; protein; 610 AA.
Human PRO polypeptide #46.
US2003017541-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 610 AA.
                                                                                                 ABO09902 standard; protein; 610 AA
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26.3%;
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Best Local Similarity 26.3%;
RESULT 1201
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Best Local Similarity 26.3%;
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US2002123463-A1.
05-SEP-2002.
(GETH ) GENENTECH INC.
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US2003032115-A1.
13-FEB-2003.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                  Best Local Similarity RESULT 1198
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                                                                  Best Local Similarity RESULT 1197
                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                ABU10837 standard;
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                                           23-JAN-2003.
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RESULT 1199
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DB 6; Length 610;

44;

Query Match

Query Match

Query Match

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Query Match

Length 610;

DB 6;

Length 610;

DB 6; 44;

Length 610;

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DB /

Length 610;

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DB 44;

Length 610;

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DB 44;

Length 610

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DB 44;

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Length 610

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DB 44;

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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                    ABM77286 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054479-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human secreted polypeptide PRO872, SEQ ID NO:92. US2003552-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104539-A1.
                                                                                                                                                                                                           Human secreted/transmembrane protein (PRO) #46.
US200306685-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA77844 standard; protein; 610 AA.
Human secreted/transmembrane.protein (PRO) #46.
US2003073180-A1.
                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein (PRO) #46 US2003068725-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted/transmembrane protein (PRO) #46 US2003068682-A1.
                                                                                                                                                                           Score 82;
Pred. No. 4
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                              610
                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 7.0%;
Local Similarity 26.3%;
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                                      Query Match 7.0%;
Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
                                                                                                                                                                                                                                                                                                                                                                protein;
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Local Similarity 26.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO35837 standard; protein
Human PRO polypeptide #46.
US2003068701-A1.
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US2003068755-Al.
10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                              ABO31515 standard;
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                                                                            RESULT 1216
                                                                                                                                                                                                                                                                                                                                               RESULT 1218
                                                                                                       Score 82; DB 6; Length 610; Pred. No. 44;
                                                                                                                                                                                                                                           Score 82; DB 6; Length 610;
Pred. No. 44;
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Pred. No. 44;
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44;
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44;
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Human secreted and transmembrane polypeptide PRO872
US2002197615-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054455-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR73973 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003036135-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR80862 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049741-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABRB1167 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049743-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM00863 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR88465 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003068743-A1.
                                    ABO08377 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044922-A1.
                                                                                                                                                                                                                                                                                                                                                                                                           ABO05584 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US200303118-A1.
13-FEB-2003.
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44;
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44;
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44;
     26.3%; Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%; Pred. No.
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RESULT 1214
                                                                                                       7.0%;
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Human PRO872 polypeptide.
US2003017981-A1.
23-JAN-2003.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1215
ID ABR89465 RF
DE Humar
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RESULT 1210
ID ABR713973 standard; pr
DE Human secreted polype
PN US2003035435-A1.
PD 20-FEB-2003.
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                                                                                                                   Best Local Similarity RESULT 1207
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Best Local Similarity
RESULT 1211
 Best Local Similarity
RESULT 1206
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                                                                                      06-MAR-2003.
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Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
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44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064452-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068705-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABM77591 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054473-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABR95260 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                                                                                                                                                                                                                                                                                                                                                  ABR94955 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92
US2003040075-A1.
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                                                               Human secreted/transmembrane protein (PRO) #46
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44;
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                                                                                                                                                                                                                                                                                                                                                    7.0%; Score 82; DB 26.3%; Pred. No. 44;
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Human secreted/transmembrane protein (PRO)
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                                                                                                             Score 82;
Pred. No. 4
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Pred. No.
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86.3%; Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                         7.0%; Score 82; 26.3%; Pred. No.
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86.3%; Pred. No.
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                                                 protein; 610 AA
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1234
ID ABM27821 standard, r
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RESULT 1230
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RESULT 1231
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             Best Local Similarity
RESULT 1225
                                                                                                                          Best Local Similarity RESULT 1226
                                                                                                                                                                                                                                        Best Local Similarity
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                                                 ABO03039 standard;
                                                                                                                                                                                                                                                     RESULT 1227
ID ABM17209 standard;
                                                                                US2003036131-A1.
20-FEB-2003.
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DB 6; Length 610;
44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064447-A1.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064440-A1.
03-APR-2003.
                                                                            Human secreted polypeptide PRO872, SEQ ID NO:92. US200306874-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                               ABM26296 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104549-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABRO2820 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064462-A1.
                                                                                                                                                                                                  ABM03608 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068722-A1.
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Human secreted/transmembrane protein (PRO) #46.
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US2003049749-Al.
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44;
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44;
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Human secreted/transmembrane protein PRO872.
US2003008297-A1.
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44;
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Pred. No.
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Pred. No.
                                                                                                                                                           Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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17-APR-2003.
(GETH ) GENENTECH INC...
7.0%;
                                             7.0%;
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                                                                                                                                                                                                                                                                    Best Local Similarity 26.3%;
RESULT 1237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 7.0%;
Local Similarity .26.3%;
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Best Local Similarity 26.3%;
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                                                                                                                                                                                                                                            10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                            (GETH ) GENENTECH INC.
                                                                                                                                                                  Best Local Similarity
RESULT 1236
                                              Best Local Similarity
RESULT 1235
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 1238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                 ABM35059 standard;
                                                                                                                                  10-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-JUN-2003
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                                                                                                                                                             Query Match
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                                           Query Match
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Length 610;

Length 610;

DB 6; 44;

6; Length 610;

DB 44;

Length 610;

9

DB 44;

Length 610;

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DB 44;

6; Length 610;

DB 44;

DB 6; Length 610; 44;

A D S E I

DB 6; Length 610;

Length 610

9

DB 44;

Query Match

P P P P P

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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                       ABM76371 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003082717-A1.
                                                                                                                                                                                                                                                                                                                         ABM76067 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104548-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM25686 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104542-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM25991 standard; protein; 610 AA.
Human secreted polypeptide PROB72, SEQ ID NO:92.
US2003104543-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA21110 standard; protein; 610 AA.
Human secreted/transmembrane polypeptide PRO872.
US2003054404-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR90600 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO03344 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003036127-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO02429 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003040061-A1.
27-FBB-2003.
                      Score 82; DB 6;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                              Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                          Score 82;
Pred. No.
                                                                                                                                                                                                                                                                            Score 82;
Pred. No.
                                                                        protein; 610 AA
                                                                                                                                                                                                                                                                          7.0%;
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                                      26.3%;
                                                                   ABO43671 standard; protein
Human PRO polypeptide #46.
US2003068732-A1.
      GENENTECH INC.
                                                                                                                         10-APR-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                                      Best Local Similarity
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                    Query Match
Best Local Similarity
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RESULT 1260
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                                                                                                                                                                                                                                                          01-MAY-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1256
                                                                                                                                                                                                                                                                                                             RESULT 1255
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                                                       RESULT 1253
                                    DB 6; Length 610;
                                                                                                                                                                                                                                                                                                           Length 610;
                                                                                                                                                                      Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 6; Length 610;
44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6; Length 610;
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44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 6; Length 610, 44;
                                                                                                                                                                                                                   ABMIS989 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064463-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABM29041 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM07017 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068699-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABM21111 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068707-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABM09457 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073175-A1.
                                                                                                Human secreted polypeptide PRO872, SEQ ID NO:92 US2003073184-A1.
                                                                                                                                                                                                                                                                                                       DB 6;
44;
                                                                                                                                                                                                                                                                                                                                                    ABO27550 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003064451-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO41327 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068695-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB
44;
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44;
                                                                                                                                                                                                                                                                                                       Score 82;
Pred. No.
                                Query Match 7.0%; Score 82;
Best Local Similarity 26.3%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 82;
Pred. No:
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                 ABM02693 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO36142 standard; protein; 610 AA
                                                                                                                                                                    7.0%;
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PA (GETH) GENENTECH INC.
QUETY MAtch 7.0%;
Best Local Similarity 26.3%;
RESULT 1249
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RESULT 1250
                                                                                                                                                                                                                                                                                                      7.0%;
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RESULT 1251
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
                                                                                                                                   17-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                       03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                    Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1246
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RESULT 1252
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RESULT 1247

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6; Length 610;
                                                                                                                                                                                                                                                                                         7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                           7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 6; Length 610;
44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 6; Length 610; 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABR71228 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003059880-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR93125 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064465-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT.1268
ID ABR93430 standard; protein; 610 AA.
DE Human secreted polypeptide PRO872, SEQ ID NO:92.
PN US2003054478-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABRR 7855 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068718-A1.
                                                                                                                                                                                                               ABR94345 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003044917-A1.
                                                                                                                                                                                                                                                                                                                                            ABR75852 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003044929-A1.
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ID ABO27855 standard; protein; 610 AA.

DE Human secreted/transmembrane protein (PRO) #46.

PN US2003064454-A1.
                                                                     ABO16920 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003054470-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA10097 standard; protein; 610 AA.
Human secreted/transmembrane protein, PRO872.
US2003059831-A1.
                                                                                                                                                            Score 82; DB
Pred. No. 44;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7.0%; Score 82; 26.3%; Pred. No.
                                                                                                                                                            7.0%;
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                  7.0%;
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Best Local Similarity 26.3%;
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RESULT 1272
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                            Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1266
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                           Best Local Similarity
RESULT 1263
                                                                                                                                                                                                                                                                         06-MAR-2003.
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                Query Match
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DB 6; Length 610; 44;
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44;
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Pred. No. 44;
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Pred. No. 44;
                                                                                  Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           6; Length 610;
                                                                                                                                                                                                                                                               ABM04887 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068727-A1.
                                                                                                                                                                                                                                                                                                                                                                                        ADMUNEAT standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003068772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABMI0372 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003069407-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABM11897 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104555-A1.
ABO29990 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003064461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABO36447 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068714-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO39497 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068776-A1.
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44;
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9
                                                                                  DB
44;
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44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                   7.0%; Score 82;
26.3%; Pred. No.
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Pred. No.
                                                                                  Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                   protein; 610 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO52043 standard; protein; 610 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 26.3%;
RESULT 1276
                                                                                                                                                                                                             Best Local Similarity 26.3%;
                                                PD 03-APR-2003.

PA (GETH) GENENTECH INC.

Query Match 7.0%;

Best Local Similarity 26.3%;

RESULT 1273
                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 26.3%; RESULT 1275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7.08;
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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RESULT 1277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #46.
US2003068758-A1.
                                                                                                                             ABO33199 standard, proteir
Human PRO polypeptide #46.
US2003068724-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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6; Length 610;

DB (

6; Length 610;

DB /

Length 610;

9

DB (

Length 610;

DB 6;

Length 610;

DB 6;

Length 610;

DB 6; 44;

Length 610;

DB 6;

Length 610;

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7.0%; Score 82; DB 6; Length 610; (6.3%; Pred. No. 44;
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Pred. No. 44;
                                                            ABM06407 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068709-A1.
                                                                                                                                                                                     ABM04218 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068716-A1.
                                                                                                                                                                                                                                                                                                              ABM22331 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068740-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM07627 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003068751-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003087374-A1.
08-MAZ-2003.
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Human secreted/transmembrane protein (PRO) #46.
US2003068684-A1.
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Pred. No.
                                                                                                                                           Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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ABOS2653 standard; protein; 610 AA
Human PRO polypeptide #46.
US2003049773-AL
                                                                                                                                       7.0%;
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                                  26.3%;
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ABM35364 standard, protein;
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(GETH ) GENENTECH INC.
    (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                              Best Local Similarity RESULT 1291
                                                                                                             10-APR-2003
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                 Query Match
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RESULT 1292
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                                                               Length 610;
                                                                                                                                                                                                                                                                                              Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Length 610;
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Pred. No. 44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054481-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR86940 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049778-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054476-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABMIS2S2 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068692-A1.
                                                             .
9
                                                                                                                                                                                                                                ABO23666 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO32125 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068733-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 6
                                                             Score 82; DB
Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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Pred. No. 4
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Pred No.
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Pred. No.
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                                                                                               Human PRO polypeptide #46.
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                                                            7.0%;
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RESULT 1288
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RESULT 1290
Human PRO polypeptide #46.
US2003049768-A1.
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US2003054987-A1.
                            13-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
RESULT 1286
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                                                                       Best Local Similarity
RESULT 1282
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Best Local Similarity
RESULT 1283
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Best Local Similarity
RESULT 1284
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Best Local Similarity
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                                                             Match
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RESULT 1287

DB 6; Length 610; 44;

Length 610;

DB 6; 44;

Length 610;

DB 6;

Length 610;

DB 6;

Length 610;

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DB 6

Length 610;

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DB 44;

Length 610;

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DB 44;

Length 610;

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DB 6 44;

Query Match

Query Match

Length 610

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DB 44;

us-10-692-730-2.spdi.rag

Query Match

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Score 82; DB 6; Length 610;
Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted polypeptide PRO872, SEQ ID NO:92.
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                                                    ABM20806 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068711-A1.
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US2003068773-A1.
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                                                                                                                                                                                                                                                                                             human secreted/transmembrane protein (PRO) #46.
US2003951-A1.
                                                                                                                                                                                  ABO41937 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049745-A1.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                 Score 82;
Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 610 AA
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           Best Local Similarity 26.3%;
RESULT 1310
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                           6; Length 610;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6; Length 610;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073169-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM13422 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064457-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABR80557 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049740-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM01168 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049770-A1.
                                                                                                                                                                                                                                                                                                                                                    ABMI8429 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054480-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR97457 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003059885-A1.
                                                                                                                                   ABO04259 standard; protein; 610 AA.

Human secreted/transmembrane protein (FRO) #46.

US2003036164-A1.

20-FB-2003.

7.0%; Score 82; DB 6;
                                                                                                                                                                                                               Best Local Similarity 26.3%; Fred. No. 44; RESULT 1302
ID ABOOSB89 standard; protein; 610 AA.
DE Human secreted/transmembrane protein (PRO) #46.
PN US2003040074-A1.
                                 ABU99207 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
                                                                                   DB /
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44;
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Human secreted/transmembrane protein PRO872
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44;
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44;
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44:
 26.3%; Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Best Local Similarity 26.3%;
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                               ABU99207 standard; protein;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1305
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RESULT 1303
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RESULT 1301
Best Local Similarity
RESULT 1300
                                                             US2003040055-A1.
27-FEB-2003.
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Query Match
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Query Match

DB 6; Length 610;

Length 610;

Length 610;

DB 6; 44;

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Score 82; DB 6; Length 610;
Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003082715-A1.
                                                                                                                                                                                                                                                                                                                                                                     ABM28736 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003082716-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM66180 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003068737-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003096359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABM34347 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003100061-A1.
                                                                                                     AB037667 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068756-A1.
                                                                                                                                                                                           Score 82; DB 6;
Pred. No. 44;
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US2003054454-A1.
20-MAR-2003.
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                                                     Score 82; DB
Pred. No. 44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 610 AA
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26.3%;
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                     10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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     US2003068742-A1.
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                                                                                                                                                                                          Query Match 7.0%; Score 82; DB 6; Length 610; Best Local Similarity 26.3%; Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                   DB 6; Length 610; 44;
                                                                     Length 610;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 6; Length 610; 44;
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44;
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Pred. No. 44;
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ABR96237 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003054458-A1.
                                                                                                                    ABM02388 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003059886-A1.
                                                                                                                                                                                                                                                          ARR65310 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2001049758-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABMI6599 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064448-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM29651 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064456-A1.
                                                                                                                                                                                                                                                                                                                                                                                                    ABR86635 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003049772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM23856 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068735-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92
US2003068753-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
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9
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44;
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Human secreted/transmembrane protein (PRO)
US2003068693-A1.
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                                                                     Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%; Pred. No.
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RESULT 1322
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1324
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RESULT 1325
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                                                                                    Best Local Similarity
RESULT 1320
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Query

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Length 610;

DB 6; 44;

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DB 6; 44;

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DB 44;

Length 610;

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DB 6; Length 610; 44;

Length 610;

DB 6; 44;

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7.0%;
            A (GETH) GENENTECH INC.
Query Match
Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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(GETH ) GENI
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Pred. No. 44;
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44;
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44;
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                                                                                                   6; Length 610;
                                                                                                                                                                                                                                                                                                                                                       6; Length 610;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068736-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM29346 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068679-A1.
                                                                                                                                                                                                                                                                                                                                                                                                    ABR85720 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049753-A1.
                                                                                                                                                ABR96542 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054460-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABM00558 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003073172-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABR99702 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
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Human secreted polypeptide PRO872, SEQ ID NO:92
                 ABO22108 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003054477-A1.
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                                                                                                                                                                                                                                                                                  ADA94329 standard; protein; 610 AA.
Human secreted/transmembrane protein PRO872
US2003059832-A1.
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44;
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44;
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Pred. No.
                                                                                                   Score 82;
Pred. No.
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                                                                 20-MAR-2003.
(GETH ) GENENTECH INC.
Query Match
7.0%;
Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity ARSULT 1347
ID ABM29346 8FF
PN Human
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RESULT 1344
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17-APR-2003.
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RESULT 1338

ID ABO22100

DE Human se

PN US200300

PD 20-MAR-:

PA (GETH )
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Score 82; DB 6; Length 610;
Pred. No. 44;
DB 6; Length 610; 44;
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44;
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44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104557-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  ABO16615 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003027276-A1.
                                                                                                                                                                                                                                                                                                                                      ADA81363 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003092121-A1.
                         Human secreted/transmembrane protein (PRO) #46. US2003068767-A1.
                                                                                                 DB 6;
44;
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US2003044920-A1.
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44;
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44;
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
 Score 82;
Pred. No.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                    Score 82;
Pred. No.
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US2003027265-Al.
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1022003054461-A1.
                                                                                                                              Human PRO polypeptide #46. US2003073182-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein; 610 AA
                                                                                                                                                                                                                                                                                       GETH ) GENENTECH INC.

Query Match
7.0%;
Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
RESULT 1352
                                                                                                 7.0%;
Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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US2003068719-A1.

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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
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                                                                                          Length 610;
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                                                                                                                                        ABM77896 standard; protein; 610 AA.

Human secreted polypeptide PRO872, SEQ ID NO:92.

US2003049783-A1.

13-MAR-2003.

A (GFH) GENENTECH INC.

7.0%; Score 82; DB 6; L
ABR81472 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                                                                                                                                                                                                                                       ABR89685 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073171-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                 ABM26601 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID.NO:92.
US2003032121-A1.
13-FEB-2003.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068702-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM03913 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068734-A1.
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                                                                                      7.0%; Score 82; DB 6;
26.3%; Pred. No. 44;
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Human secreted/transmembrane protein (PRO) #46.
US2003064464-A1.
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Human secreted/transmembrane protein (PRO) #46.
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Pred. No. 44;
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Pred. No.
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                             US2003049744-A1.
113-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1359
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RESULT 1361
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RESULT 1362
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RESULT 1358
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RESULT 1364
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Best Local Similarity
RESULT 1366
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Score 82; DB 6; Length 610;
Pred. No. 44;
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44;
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Pred. No. 44;
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Human secreted/transmembrane protein (PRO) #46.
US2003049750-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABO50518 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049779-A1.
13-MAR-2003.
                                                             ABO41632 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
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44;
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Human secreted/transmembrane protein (PRO)
US2003049747-A1.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Human PRO polypeptide #46.
US2003068738-A1.
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Human PRO polypeptide #46.
US2003049766-A1.
                       7.0%;
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                                                                                                                     Best Local Similarity 26.3%;
RESULT 1368
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                                                            protein;
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Local Similarity 26.3%;
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10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 1369
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Best Local Similarity
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Best Local Similarity
RESULT 1373
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Best Local Similarity
RESULT 1374
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                                Best Local Similarity RESULT 1367
                                                                                                                                                                                                                                                                                                                                Best Local Similarity
                                                            ABO41632 standard;
                                                                                       US2003068729-A1.
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                        Query Match
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RESULT 1372
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RESULT 1371
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13-MAR-2003. (GETH ) GENENTECH INC. US2003068696-A1. 17-APR-2003. Query Match Query Match 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44; 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44; 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44; Score 82; DB 6; Length 610; Pred. No. 44; DB 6; Length 610; 44; 7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44; Score 82; DB 6; Length 610; Pred. No. 44; DB 6; Length 610; 44; DB 6; Length 610; 44; 6; Length 610 ABMI6904 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040078-A1. ABR96847 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003054462-A1. ABM12202 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003064445-A1. ABR79642 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003040059-A1. ABM16294 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003064449-A1. ABM24161 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. US2003064441-A1. ABM14642 standard; protein; 610 AA. Human secreted polypeptide PRO872, SEQ ID NO:92. ABO17936 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003044918-A1. ABO20888 standard; protein; 610 AA. Human secreted/transmembrane protein (PRO) #46. US2003032132-A1. ADA38554 standard; protein; 610 AA. Human secreted/transmembrane protein PRO872. US2003059780-A1. 7.0%; Score 82; DB 26.3%; Pred. No. 44; Score 82; Pred. No. 4 Score 82; Pred. No. Score 82; Pred. No. 7.0%; 7.0%; Best Local Similarity 26.3%; RESULT 1381 7.0%; 26.3%; 7.0%; (GETH ) GENENTECH INC. Best Local Similarity RESULT 1383 Best Local Similarity RESULT 1384 Best Local Similarity RESULT 1378 Best Local Similarity RESULT 1380 Best Local Similarity RESULT 1382 Best Local Similarity Best Local Similarity Best Local Similarity 06-MAR-2003. 13-FEB-2003. 03-APR-2003 03-APR-2003 27-FEB-2003 20-MAR-2003 Query Match RESULT 1376 RESULT 1379 RESULT 1385

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DB 6; Length 610;
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                                                  Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104545-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104541-Al.
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44;
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44;
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44;
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44;
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                       Score 82;
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Pred. No.
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                     Best Local Similarity 26.3%;
RESULT 1386
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RESULT 1388
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Best Local Similarity 26.3%;
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Human PRO polypeptide #46.
US2003049765-Al.
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US2003049762-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
10-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                                T 1386
ABM04523 standard;
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Query Match

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7.0%; Score 82; DB 6; Length 610; 26.3%; Pred. No. 44;
                                                                                                                DB 6; Length 610; 44;
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Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032133-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003032136-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
บุนรถ03040057-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054467-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US20034469-A1..
20-MAR-2003.
(GETH ) GENENTECH INC.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003022770-Al.
06-FEB-2003.
                                       ADA83161 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049752-A1.
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Human secreted/transmembrane protein (PRO) #46.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
Best Local Similarity
RESULT 1395
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Best Local Similarity
RESULT 1401
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RESULT 1404
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Best Local Similarity
RESULT 1403
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Score 82; DB 6; Length 610;
Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003059879-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049738-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064469-A1.
                                            ABO20583 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US203032126-A1.
13-FEB-2003.
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Human secreted/transmembrane protein (PRO) #46.
US2003064466-A1.
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Human secreted/transmembrane protein PRO872.
US2003060407-A1.
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Pred. No. 44;
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Pred. No. 4
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Pred. No.
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Human PRO polypeptide #46.
US2003054466-A1.
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Human PRO polypeptide #46.
US2003054463-A1.
20-MAR-2003.
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Human PRO polypeptide #46.
US2003064453-A1.
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ABO30905 standard; protein; 610 AA.
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Best Local Similarity
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RESULT 1405
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10-APR-2003. (GETH ) GENENTECH INC.

US2003068688-A1.

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Score 82; DB 6; Length 610;
Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
                                                            Length 610;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068759-A1.
                                                                                                     ABM27211 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068760-A1.
                                                                                                                                                                                                                            ABM29956 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068769-A1.
                                                                                                                                                                                                                                                                                                                                                 ABMO5492 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003045700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABM66685 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92
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Human secreted/transmembrane protein (PRO) #46.
US2003068765-A1.
Human secreted/transmembrane protein (PRO) #46.
US2003064468-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO42242 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049748-A1.
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                                                        DB
44;
                                                           Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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10-APR-2003.
(GETH ) GENENTECH INC.
7.0%; St
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Best Local Similarity 26.3%;
RESULT 1417
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                             (GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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RESULT 1419
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RESULT 1422
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RESULT 1423
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RESULT 1421
                                                                         Best Local Similarity
                                03-APR-2003
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                                                              Query Match
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                                                                                         RESULT
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Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                      Length 610;
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                                                                                                                                    Human secreted polypeptide PRO872, SEQ ID NO:92.052003104552-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABM26906 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nemussus standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068763-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABR88160 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068720-A1.
                        Human secreted/transmembrane protein (PRO) #46.0122003082767-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AB039802 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068689-A1.
                                                                                                                                                                                                                                                                                                                                                                                     ABO49603 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049775-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA78456 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003073181-A1.
                                                                                                      DB 6;
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44;
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44;
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Pred. No.
                                                                                                      Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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26.3%; Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                     protein; 610 AA
7.08;
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(GETH ) GENENTECH INC.
          Best Local Similarity 26.3%;
RESULT 1424
                                                                                                                                                                                                                                                                                                                                          Watch 7.0%;
Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
RESULT 1428
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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                                                                                                                                                                                                                                                                                                             13-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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   Query Match
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RESULT 1434

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03-APR-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                     DB 7; Length 610;
               Score 82; DB 6; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                          Score 82; DB 7; Length 610;
Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABM17819 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040072-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABR95870 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040073-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABR77062 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003044927-A1.
                                                                ABO49908 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049776-A1.
                                                                                                                                                                                                                                                                                                                        Human secreted/transmembrane protein (PRO) #46.
US2003036126-A1.
20-FEB-2003.
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Human secreted/transmembrane protein (PRO) #46.
US2003054475-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO24276 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003064467-A1.
                                                                                                                                                                                                    ABOS0823 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46
US2003049780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABO19973 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003032124-A1.
                                                                                                                                                     Score 82;
Pred. No.
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Pred. No.
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
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               Query Match
Best Local Similarity
RESULT 1433
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Best Local Similarity
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Best Local Similarity
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RESULT 1441
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RESULT 1439

RESULT 1442

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DB 7; Length 610; 44;
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                                                                                                                Length 610;
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44;
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44;
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                                        ABR86025 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003049759-A1.
                                                                                                                                                       ABM10677 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064455-A1.
                                                                                                                                                                                                                                                              Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104550-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted polypeptide PRO872, SEQ ID NO:92 US2003068764-A1.
                                                                                                                DB 7;
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44;
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44;
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                                                                                                               Score 82;
Pred. No. 4
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Pred. No. 4
  Score 82;
Pred. No.
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Pred. No.
                                                                                                                                                                                                                               Score 82;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein, 610 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                             7.0%;
                                                                                                                                                                                                                              7.0%;
Local Similarity 26.3%;
Query Match 7.0%;
Best Local Similarity 26.3%;
                                                                                                               y Match
Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Local Similarity 26.3%;
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Local Similarity 26.3%;
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US2003068728-Al.
                                                                                   13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                               20-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                03-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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US2003059882-A1.
                                                                                                                                                                                                                                                                              DB 7; Length 610;
7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
                                                                                                                                  Score 82; DB 7; Length 610;
Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                7; Length 610;
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44;
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Pred. No. 44;
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036161-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABR89075 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036119-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR72448 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036120-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABR80252 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003049739-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM01473 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
                                            ABM19281 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104551-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR69036 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003027273-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO18546 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003044921-A1.
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44;
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Human secreted/transmembrane protein (PRO)
US2003049757-A1.
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26.3%; Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Human PRO polypeptide #46.
US2003049761-Al.
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26.3%;
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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RESULT 1460
ID ABR80025 standard; pr
DE Human secreted polype
PD 13-MAR-2003
PA (GETH ) GENENTECH INC
Query Match
Best Local Similarity
RESULT 1452
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RESULT 1459
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Best Local Similarity
RESULT 1456
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RESULT 1458
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RESULT 1461
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DB 7; Length 610; 44;
                              DB 7; Length 610;
44;
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44;
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                                                               Human secreted polypeptide PRO872, SEQ ID NO:92.07.2003059884-A1.
                                                                                                                                                                                                ABR87245 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068687-A1.
                                                                                                                                                                                                                                                                                                             Human secreted polypeptide PRO872, SEQ ID NO:92.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064443-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064444-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068686-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABM09762 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073178-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO29380 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068697-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO11210 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068710-A1.
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44;
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44;
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44;
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Pred. No.
                              Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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RESULT 1469
                         Best Local Similarity 26.3%;
RESULT 1462
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RESULT 1464
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RESULT 1467
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Local Similarity 26.3%;
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Best Local Similarity 26.3%;
RESULT 1465
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Best Local Similarity 26.3%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
27-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                       (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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20-MAR-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
                                                                                                  ABM01778 standard;
     US2003054457-A1.
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                                                   Query Match
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                  vuery Match 7.0%; Score 82; DB 7; Length 610; Best Local Similarity 26.3%; Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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Pred. No. 44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABR77667 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040067-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABR78877 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003054456-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
                                                                       ABO38887 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068774-A1.
10-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                               Human secreted/transmembrane protein (PRO) #46. US2003049781-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein (PRO) #46. US2003036158-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 82; DB 7;
Pred. No. 44;
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Human secreted/transmembrane protein (PRO)
US2003054482-A1.
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Pred. No.
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US200330451-A1.
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17-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                         Best Local Similarity
RESULT 1472
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Best Local Similarity
RESULT 1479
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Best Local Similarity
RESULT 1480
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D N C

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Length 610;
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    Length 610;
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                                                                                                                                                            Human secreted polypeptide PRO872, SEQ ID NO:92. US2003049764-Al.
                                                               Human secreted polypeptide PRO872, SEQ ID NO:92 US2003059883-A1.
                                                                                                                                                                                                                                                                                              ABR89990 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003073177-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                   ADA22236 standard; protein; 610 AA.
Human secreted/transmembrane polypeptide PRO872
US2003040473-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABM27516 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92
US2003064442-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABM13117 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003064450-A1.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068683-A1.
10-APR-2003.
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Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003068754-A1.
   DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO31820 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068731-A1.
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44;
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44;
                                                                                                                           DB 7;
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Pred. No. 44;
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Pred. No. 44;
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44;
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Pred. No. 4
   Score 82;
Pred. No.
                                                                                                                           Score 82;
Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
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Pred. No.
7.0%;
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26.3%;
                                                                                                                         Match 7.0%;
Local Similarity 26.3%;
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Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
RESULT 1488
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                 Best Local Similarity 26.3%;
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Best Local Similarity 26.3%;
                                                 protein;
                                                                                                                                                                                                                   13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                          17-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                         (GETH ) GENENTECH INC.
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Best Local Similarity
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us-10-692-730-2.spdi.rag

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7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
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7.0%; Score 82; DB 7; Length 610; 26.3%; Pred. No. 44;
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Pred. No. 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABM20196 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003104556-A1.
                                                                                                                                                                                                        ABM74542 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003096351-A1.
                                                                                                                                                                                                                                                                                                                                                                    ABM313737 standard; protein; 610 AA.
Humman secreted polypeptide PRO872, SEQ ID NO:92.
US2003096358-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABR72753 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003036122-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABR85110 standard; protein; 610 AA.
Human secreted polypeptide PRO872, SEQ ID NO:92.
US2003040065-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO48688 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003049756-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO15395 standard; protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003036121-A1.
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Human secreted/transmembrane protein (PRO) #46.
                                                          ABO40107 standard, protein; 610 AA.
Human secreted/transmembrane protein (PRO) #46.
US2003068681-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AB022498 standard, protein, 610 AA.
Human secreted/transmembrane protein PRO872.
US2003017982-A1.
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20-FEB-2003.
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Best Local Similarity
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RESULT 1492
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RESULT 1493
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Best Local Similarity
RESULT 1496
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               Best Local Similarity
RESULT 1490
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RESULT 1499
ID ABO15090 standard;
DE Human secreted/tran
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Score 82; DB 7; Length 610; Pred. No. 44;
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Pred. No. 44;
PN US2003044919-A1.

PD 06-MAR-2003.

Query Match
Best Local Similarity 26.3%; Pred. No. 44;

RESULT 1500

ID AB017225 standard; protein; 610 AA.

DE Human secreted/transmembrane protein (PRO) #46.

PN US2003040077-A1.

PD 27-FRB-2003.

Query Match

7.0%; Score 82; DB 7; Lea
                                                                                                                                                                                                 7.0%;
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Length 229;
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(GETH ) GENENTECH INC.
1747 Match 76.0%; Score 174; DB 3; Length 229; Ft Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human PRO733 (UNQ411) protein sequence SEQ ID NO:447.
WO200053756-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB34734 standard; protein; 229 AA.
Human secreted protein encoded by DNA clone vq20 1.
WO200055375-A1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USACOLIO.
29-JUL-2004.
29-JUL-2004.
100.08; Score 229; DB B; I
(ery Match 100.08; Pred. No. 1.5e-211;
                                                                                                                                                                                                                    PD 23-OCT-2003.

PA (HUMA-) HUMAN GENOME SCI INC.

Querry Match 100.0%; Score 229; DB 8; I

Best Local Similarity 100.0%; Pred. No. 1.5e-211;

RESULT 8
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Best Local Similarity 100.0%; Pred. No. 1.5e-211;
                                                                                                    100.0%; Score 229; DB 7; 100.0%; Pred. No. 1.5e-211;
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WO9836068-A2.
                                                                                                                                                                                                                                                                                                                ADL91839 standard; protein; 229 AA.
Human PRO83480 protein SEQ ID NO:60
WO2004024076-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human TIR-like ligand II protein. US2004146501-A1.
                                                                                                                                                              protein; 229 AA
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Human protein clone HP02515.
WO200005367-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW49100 standard; protein; 229 AA
                                                                                                                                                                                  Human T1R-like ligand II protein. US2003198618-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY41754 standard; protein; 22
Human PRO733 protein sequence.
WO9946281-A2.
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(SAGA ) SAGAMI CHEM RES CENT.
(PROT-) PROTEGENE INC.
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(ALPH-) ALPHAGENE INC.
76.0%;
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(INCY-) INCYTE PHARM INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                      (NIJJ/) NI J.
(GENTZ R.
(RUBE/) RUBEN S M.
                                                                                                                      Local Similarity
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        US2003100048-A1.
                            29-MAY-2003
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                                          (without alignments)
541.340 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of SUMMARIES
                                                                                                                    MGDKIWLPFPVLLLAALPPV.....SAIQVYMLKSLFEDKRKSRT 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Description
OM protein - protein search, using sw model
Run on: March 19, 2007, 23:13:05 ; Search time 207 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAE02806 standard; protein; 229 AA.

Human TIR-like ligand II protein.

W0200142506-AI.

14-JUN-2001.

14-JUNA-) HUMAN GENOME SCI INC.

ery Match

100.0%; Score 229; DB 4; Length 229; St Local Similarity 100.0%; Pred. No. 1.5e-211;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; SCI INC.
100.0%; Score 229; DB 2; Length 229;
100.0%; Pred. No. 1.5e-211;
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(EOSB-) EOS BIOTECHNOLOGY INC.
ery Match
ery Match
incomplexity 100.0%; Pred. No. 1.5e-211;
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ry Match 100.0%; Score 229; DB 7; Length 229;
t Local Similarity 100.0%; Pred. No. 1.5e-211;
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Angiogenesis protein AAA4 protein sequence (Fig 4).
WO200111086-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human T1 receptor (T1R)-like ligand II polypeptide
US6586210-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 229; DB 6; 100.0%; Pred. No. 1.5e-211;
                                                                                                                                                    Gapop 60.0 , Gapext 60.0 2782104 seqs, 489333398 residues Total number of hits satisfying chosen parameters: Minimum DB seq length: 2000000000
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Human T1-receptor-(T1R-)like ligand II.
US2002142461-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW48335 standard; protein; 229 AA.
Human T1 receptor-like ligand II.
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Human T1 receptor-like ligand II.
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J.GEVTZ R L

J.GEV ) RUBEN S M.

J.Y MAtch

Best Local Similarity

RESULT 5

ID ADB87500 stand

DE Human T1 r

PN US5867

PD 01
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(HUMA-) HUMAN GENOME
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Best Local Similarity
RESULT 4
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                                                                                                  Perfect score:
                                                                                                                                      Scoring table:
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Human secreted/transmembrane protein (PRO) #205.
US2003036179-A1.
                                       20-FEB-2003
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 6; Length 229;
(ery Match 76.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2002192706-A1.
                                                                                                                                       ABO17820 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003032156-A1.
                                                                                                                                                                                                                                                                                 ABO25256 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003050239-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABUB4942 standard; protein; 229 AA.
Human secreted and transmembrane PRO polypeptide #18
US2002177553-A1.
                                                00-JUN 2001.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 4; Leery Match 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                        13-FEB-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 6; Leery Match 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
.ry Match
.roman eimilarity 100.0%; Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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Novel secreted and transmembrane protein PRO733.
US2003017563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #205.
192003036180-Al.
AAU12376 standard; protein; 229 AA.
Human PRO733 polypeptide sequence.
WO200140466-A2.
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Human PRO733 polypeptide.
US2002169284-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 229 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human.
US20030361bv
20-FEB-2003.
'TFTH ) GENENTECH INC.
76.0%; PO.
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US2003004311-A1.
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Best Local Similarity
RESULT 17

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PD 13-**
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                                                                                                     Best Local Similarity RESULT 16
                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 18
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RESULT 20
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                                                                                                                                                                                                                                                                                                                                                                                                                            ABU81074 standard;
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Length 229;
                                                                                                                                                                        76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
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(GETH) CENENTECH INC.
PRACCH 76.0%; Score 174; DB 6; Length 229;
rt Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                ABU67050 standard; protein; 229 AA.
Human secreted/transmembrane, PRO, protein SEQ ID 410.
US2003032155-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUS203022128-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB19418 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003068796-A1.
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(GETH ) GENENTECH INC.
:ry Match '76.0%; Score 174; DB 6; Le
.... etmilarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
:ry Match 76.0%; Score 174; DB 6; Le
.ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
:ry Match 76.0%; Score 174; DB 6; LK
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Pred. No. 1.2e-158;
(GETH ) GENENTECH INC.

ry Match
76.0%; Score 174; DB 6; L

r Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
                                                                                                                                                                                                                            ABG72289 standard; protein; 229 AA.
Human p24 vesicle trafficking protein Hp24-2.
US2002132999-A1.
                                                                           ABU80409 standard; protein; 229 AA.
Human secreted/transmembrane protein PRO733.
US2003004102-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA19010 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003054517-A1.
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Human PRO polypeptide #205.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA61633 standard; protein; 229 AA
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19-SEP-2002.

(INCY-) INCYTE PHARM INC.

OPT MATCh ...:larity 100.0%;
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10-APR-2003.
(GETT ) GENENTECH INC. 76.0%; S
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(GETH ) GENENTECH INC.
76.0%;
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(GETH ) GENENTECH INC.
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US2003073212-A1.
                                                                                                                                                                                    Best Local Similarity
RESULT 26
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                         Query Match
Best Local Similarity
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US2003049816-A1.
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Length 229;

Length 229;

Length 229;

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vuely match 76.0%; Score 174; DB 6; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 49
                                                                                                                                                                                                                                                                                                                                      Score 174; DB 6; Length 229;
Pred. No. 1.2e-158;
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(GET) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003082694-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel human secreted and transmembrane protein PRO733.06-MAR-2003.
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Novel human secreted and transmembrane protein PRO733
US2003050241-A1.
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Novel human secreted and transmembrane protein PRO733
US2003073211-A1.
   08-MAY-2003.
(GETH ) GENENTECH INC.
1ry Match 76.0%; Score 174; DB 6; Le
1ry Match 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 6; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                           ADB16743 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087349-A1.
                                                                                                                                                                                                                                                                                                                                                                                          ADB14898 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003087351-A1.
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192003077722-A1.
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US200082710-A1.
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08-MAY-2003.
(GETH ) GENENTECH INC.
76.0%; SC
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Query Match
76.0%; 8
Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                   Best Local Similarity
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Best Local Similarity
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                                      Query Match
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(GETH) GENENTECH INC.
(EFTH) GENENTECH INC.
FLY Match 76.0%; Score 174; DB 6; Length 229;
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GENENTECH INC.
ry Match
ry act. 100.0%; Score 174; DB 6; Length 229;
rocal Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(SETH ) GENENTECH INC.
ery Match
76.0%; Score 174; DB 6; Length 229;
                                                       DB 6; Length 229;
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                                                                                             ALMANASH Standard; protein; 229 AA.

Novel human secreted and transmembrane protein PRO733.
US203082111-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003082693-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA87541 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087345-A1.
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10-APR-2003.
(GRTH) GENENTECH INC.
6ry Match 76.0%; Score 174; DB 6; Le
ery Match 76.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
ery Match 76.0%; Score 174; DB 6; Leery Match 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.

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16.0%; Score 174; DB 6; Le

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100.0%; Pred. No. 1.2e-158;
                                                     76.0%; Score 174; DB 6; L4 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 6; LK
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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US2003087350-A1.
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17-App.
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Human PRO polypeptide #205.
US2003068795-A1.
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Human PRO polypeptide #205.
US2003068794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA97098 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003082705-A1.
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US2003082763-A1.
                 01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 40
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                                                                Best Local Similarity
RESULT 34
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Best Local Similarity
RESULT 39
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US2003082704-A1.
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                                                       Ouery Match
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PD 24-APR-2003.

PA (GETH) GENENTECH INC.

Query Match 76.0%; Score 174; DB 6; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 65
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(GETH) GENENTECH INC.

(GY) Match

76.0%; Score 174; DB 6; Length 229;

Fred, No. 1.2e-158;
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(GETH ) GENENTECH INC.

(ery Match 76.0%; Score 174; DB 6; Length 229; ery Match 100.0%; Pred. No. 1.2e-158;
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BE Human PRO polypeptide #205.

PN US2003082703-A1.

PD 01-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match

76.0%; Score 174; DB 6; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
Query Match 76.0%; Score 174; DB 6; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 61
                                                                                                                                                                                                       (GETH) GENENTECH INC.

ry Match 76.0%; Score 174; DB 6; Length 229; t Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CULLY MATCH 76.0%; Score 174; DB 6; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CATANA GENENTECH INC. 76.0%; Score 174; DB 6; Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76.0%; Score 174; DB 6; L
100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide SEQ ID NO 410. US2003077715-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO polypeptide SEQ ID NO 410.
US2003077714-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #205.
15-May An.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #205.
US2003073210-A1.
                                                                                               ADA80566 standard, protein; 229 AA. Human PRO polypeptide #205.
US2003082761-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA93505 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003077721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA61070 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USACOCO
17-APR-2003.
(GETH ) GENENTECH INC.
76.0%;
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Human PRO polypeptide #205.
US2003096386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 67
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Best Local Similarity
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US2003049817-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA84782 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082708-A1.
US2003082708-A1.
(GETH ) GENENTECH INC.
ery Match
st Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                             76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
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                                            Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003082695-A1.
                                                                                                                        ABO19711 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003050240-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   USEVEL.
01-MAY-2003.
(GETH ) GENENTECH INC.
1ery Match 76.0%; Score 174; DB 6; Let
1ery Match 100.0%; Pred. No. 1.2e-158;
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USACCOURTS
(GETH) GENENTECH INC.
(GETY Match 76.0%; Score 174; DB 6; Lf.
(ery Match 74.0%; Pred. No. 1.2e-158;
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Ocarda (GETA) GENENTECH INC.
(GETA) GENENTECH INC.
76.0%; Score 174; DB 6; LA
ery Match
` ``milarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PD 24-APR-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                            76.0%; Score 174; DB 6; L 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane polypeptide PRO733 US2003055216-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB24769 standard; protein; 229 AA. Human PRO polypeptide SEQ ID NO 410 US20030777713-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PRO polypeptide #205.
US2003082701-A1.
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USZ003073214-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA74536 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003068798-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA75256 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003073216-A1.
                                                                                                                                                                                                                                                                                                                                                 ADA12647 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LACATION CONTROL CONTR
                                                                                                                                                                                                       13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAR-2003.
(GETH ) GENENTECH INC.
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                (GETH ) GENENTECH INC
                                       Query Match
Best Local Similarity
RESULT 52
ID A8019711 standard; pr.
DE Novel human secreted
PN US2003050240-A1.
PD 13-MAR-2003
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 53
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RESULT 55

Query Match

Query Match

Query Match

Query Match

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20-MAR-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
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(GETH) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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4 (GETH) GENENTECH INC.

Query Match 76.0%; Score 174; DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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        ADA88093 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082700-A1.
                                                                                                                                                        ADA46481 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003054516-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA88645 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003073213-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB22340 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087344-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2005.
(GETH ) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 7; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 7; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 174; DB 7; Louis Pred. No. 1.2e-158;
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Beet Local Similarity '100.0%; Pred. No. 1.2e-158;
RESULT 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76.0%; Score 174; DB 7; L 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #205. US2003082706-Al.
                                                                                                                                                                                                                                                                                                                                    ADB28511 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082699-A1.
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0/02003059909-Al.
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Human PRO polypeptide #205.
US2003022239-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA67031 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003068793-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US20030..._
17-APR-2003.
(GETH ) GENENTECH INC.
"atch 76.0%; SCO
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US2003082686-A1.
                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA97650 standard;
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                                                                                                                                                    Length 229;
                                                                                                                                                                                                                                                                                                                 76.0%; Score 174; DB 6; Length 229; 100.0%; Pred. No. 1.2e-158;
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Beet Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABO19602 standard; protein; 229 AA.
Novel human secreted and transmembrane polypeptide #70
US2003049633-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB21788 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA86990 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082709-A1.
        100.0%; Pred. No. 1.2e-158;
                                                                                                                                                76.0%; Score 174; DB 6; L 100.0%; Pred. No. 1.2e-158;
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                                            ADA96546 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082690-Al.
                                                                                                                                                                                                             ADA81118 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003082702-A1.
                                                                                                                                                                                                                                                                                                                                                                        ADA95994 standard, protein; 229 AA.
Human PRO polypeptide #205.
US2003082759-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB26303 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003082760-A1.
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US200307710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA/7567 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003068797-A1.
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DE Novel human secreted ar
PD US2003082709-A1.
DD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                         01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC
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                                                                                                                                                                 Best Local Similarity RESULT 71
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Best Local Similarity RESULT 70
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Query Match

Length 229;

Length 229;

Length 229;

Length 229;

Query Match

Query Match RESULT 76

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76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                            PA (GETH) GENENTECH INC. Obery Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-2003.
(GETH) GENENTECH INC.
76.0%; Score 174; DB 7; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
127 Match 3 76.0%; Score 174; DB 7; Length 229; St Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
Query Match
76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human secreted and transmembrane protein PRO733. US2001082687-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003082696-A1.
                                                                                                                                                        ADB39535 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082764-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USACCOLOR OL MAY-2003.

(GETH, GENENTECH INC. 76.0%; Score 174; DB 7; Lety Match chilarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 100
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Human PRO polypeptide #205.
US2003082697-A1.
USA047-2003.
(GETH) GENENTECH INC.
T6.0%; Score 174; DB 7; L.

T Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide SEQ ID NO 410. V22003077716-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB34527 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077717-A1.
                                                                                                                                                                                                                                                                                                                ADB73953 standard; protein; 229 AA. Human PRO polypeptide #70. US2003045462-A1. G6-MAR-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 229 AA
                                                        (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #70.
US2003083248-Al.
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(GETH ) GENENTECH INC.
                                                                                                            Best Local Similarity RESULT 98
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Best Local Similarity
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                                                                                               Query Match
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Best Local Similarity 100.0%; Score 174; DB 7; Length 229; RESULT 96

ID ADB89702 standard; protein; 229 AA.

PN US2003082698-AI.

PA O1-MAY-200-
Luery Match Match T6.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT B TD ADB22892 standard; protein; 229 AA. DE Human PRO polypeptide #205. PN US2003077711-A1. PD 24-APR-20^2
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                                                                                                                                                                                                                                                              Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 90
                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003082712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB38702 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003082766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB38150 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087347-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAY-2003.
(GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Lu
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
76.0%; SCOTE 174; DB 7;
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Human PRO polypeptide SEQ ID NO 410.
US2003077712-A1.
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Human PRO polypeptide #205.
US2003087352-A1.
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Human PRO polypeptide #205.
US2003082762-A1.
01-MAY-2003.
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24-APR-2003.
(GETH ) GENENTECH INC.
"a+ch 76.0%; SCO
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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RESULT 92

RESULT 95 ID ADB66

Length 229;

Length 229;

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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 118
                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 119
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                      RESULT 116
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RESULT 121
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                                           BEEE
76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                               Length 229;
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PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                ADB46578 standard, protein, 229 AA.

Novel human secreted and transmembrane protein PRO733
US2003082692-A1.
                                                                                                GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

76.0%; Score 174; DB 7; Le
                                                                                                                                                                                                                                                24-APR-2203.
24-APR-2203.
(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 7; Le
(ery Match 76.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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ry Match

t Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC61855 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003049684-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC44095 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003054986-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC66919 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003060406-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC63819 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003054405-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC69043 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC63103 standard, protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003068648-A1.
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(GETH ) GENENTECH INC.
76.0%; SCOIE 174;
                                                                                                                                                                                                  ADB36183 standard; protein; 229 AA. Human PRO polypeptide SEQ ID NO 410. US2003077720-A1.
                                                 ADB35079 standard; protein; 229 AA.
Human PRO polypeptide SEQ ID NO 410.
US2003077718-A1.
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01-MAY-2003.
(GETH ) GENENTECH INC.
MAtch "Totty 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 115
           Best Local Similarity
RESULT 107
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Best Local Similarity
RESULT 109
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Best Local Similarity
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Query Match
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Length 229;
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Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein Seq ID410.
US2203087365-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003092106-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003092107-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC59977 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003092105-A1.
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(GETH) GENENTECH INC.

(GETY Match

76.0%; Score 174; DB 7; Le

rery Match

76.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 7; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
?ry Match 76.0%; Score 174; DB 7; Ld
                                                                          10-APR-2003.
(GETH) GENENTECH INC.
76.0%; Score 174; DB 7; L
BEST Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
FLY MATCh 76.0%; Score 174; DB 7; Lost Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                     76.0%; Score 174; DB 7; I 100.0%; Pred. No. 1.2e-158;
ADC68168 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003069178-A1.
                                                                                                                                                                                                ADC41488 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003072745-A1.
                                                                                                                                                                                                                                                                                                                                                                                        ADC67543 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003073131-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC62479 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003073624-A1.
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Human secreted/transmembrane protein, PRO733.
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                                                                                                                                                                                                                                                           17-APR-2003.
(GETH ) GENENTECH INC: 76.0%;
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) GENENTECH INC.
76.0%;
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ADC80043 standard; protein; 229 AA
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(GETH ) GENENTECH INC.
(SEY Match 76.0%; Score 174; DB 7; Length 229;
ery Match 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                     Length 229;
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                                                                                                                         Length 229
                  ADC57338 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seq ID410.
US2003087366-A1.
08-MAY-2003.
GETH) GENENTECH INC.
176.0%; Score 174; DB 7; Length 22 et Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADCS4629 standard; protein; 229 AA.

Novel human secreted and transmembrane protein Seg ID410.
US2003087363-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC53590 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg 1D410.
US2003087364-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADCS9113 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC55991 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seq ID410.
US2003087360-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADCS8561 standard; protein; 229 AA.
Novel human secreted and transmembrane protein Seg ID410.
US2003087346-A1.
                                                                                                                                                                               ADC60529 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                                                                                              ADC51004 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087361-A1.
                                                                                                                                                                                                                                                                               76.0%; Score 174; DB 7; Lo
100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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Human PRO polypeptide #205.
US2003087362-A1.
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100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 129
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                                                                                                                                                                                                                                                                                                  Best Local Similarity
RESULT 127
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                                                                                                                                   Best Local Similarity
RESULT 126
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                         US2003087367-A1.
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RESULT 125
1D ADC573
DE Novel
PN US2003
PD 08-MAY
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RESULT 133
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PD 15-MAY-2003.

PA (GETH) GENENTECH INC.

Query Match 76.0%; Score 174; 'DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 135
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(GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 229;
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Pred. No. 1.2e-158;
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ry Match 76.0%; Score 174; DB 7; Length 229; t Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US203092104-A1.
                                                                                                                                                                      ADC90227 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087348-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD04639 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC80595 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003092103-A1.
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Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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16.0cm 2.2.
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Human PRO polypeptide #205.
US2003194773-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD10064 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194776-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD11102 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194774-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC47983 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003194771-A1.
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16-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; S
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(GETH ) GENENTECH INC.
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GENENTECH INC.
76.0%;
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(GETH ) GENENTECH INC.
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Length 229;

Length 229;

Length 229;

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76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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1 (GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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1 (GETH) GENENTECH INC.

Query Match

96st Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE32351 standard, protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003194765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PD 23-OCT-2003.
PA (GETT) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 156
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(GETH ) GENENTECH INC.
:ry Match 76.0%; Score 174; DB 7; Le:
:ry Match 100.0%; Pred. No. 1.2e-158;
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1 (GETH) GENENTECH INC.

76.0%; Score 174; DB 7; L

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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( GETH) GENENTECH INC.

76.0%; Score 174; DB 7; L

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003203434-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE16649 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203435-A1.
                                                                                                                                        Human secreted/transmembrane protein, PRO733 US2003096744-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                     Human PRO polypeptide #205.
US2003199055-A1.
                                                                                                                                                                                                                                                                           ADD92544 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003199030-A1.
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Human PRO polypeptide #205.
US2003199057-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE22283 standard; protein; 229 AA. Human PRO polypeptide #205; US2003199056-A1.
                                                                                                                        protein; 229 AA
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30-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; SCO
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US2003203428-A1.
                         30-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                  23-OCT-2003.
(GETH ) GENENTECH INC.
                                                                               Best Local Similarity
RESULT 153
                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                        ADE49481 standard;
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       US2003203432-A1.
                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 155
                                   08-WAY-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
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Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 152
ID ADD54227 standard; protein; 229 AA.
DE Novel human secreted and transmembrane protein PRO733.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                               ADD1225 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733. US2003203438-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD53656 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003203437-A1.
                                                                                                                                                                                                                                                                                                                                                      30-OCT-zvus.
(GETH ) GENENTECH INC.
rry Match 76.0%; Score 174; DB 7; Le
'' .... Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
iry Match 76.0%; Score 174; DB 7; Le
... ran similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                   DB 7;
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(GETH ) GENEWIECH INC.
76.0%; SCOIE 174;
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                                                                                                                          Human PRO polypeptide #205.
US2003194775-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD52364 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD53104 standard; protein; 229 AA. Human RRO polypeptide #205.
1022003194792-A1.
16-0CT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD51812 standard, protein; 229 AA.
Human PRO polypeptide #205.
US2003194779-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD02611 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003203431-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD02045 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003203430-A1.
                                                                                                                                                                                                                                                     100.08;
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.2003.
.iH ) GENENTECH 1.
.ery Match
Best Local Similarity 1
RESULT 145
ID ADD41225 stander
DE Novel humar
PN US2003?
PD 30-C
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RESULT 146
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Best Local Similarity
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RESULT 149
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Best Local Similarity
RESULT 150
                                                                                            Best Local Similarity
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                     US2003087358-A1.
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                                                                                                                  RESULT 144
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PD 23-OCT-2003.

PD A (GETH ) GENENTECH INC.

Query Match

Query Match

T6.0%; Score 174; DB 7; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 172

ID ADE18964 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OSCT_2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
ery Match 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                             Vuely match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 173
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(GETH) GENENTECH INC.
76.0%; Score 174; DB 7; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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1 (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
21Y Match 76.0%; Score 174; DB 7; Length 229;
3t Local Similarity 100.0%; Pred. No. 1.2e-158;
                                  Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 171
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Novel human secreted and transmembrane protein PRO733.
US2003194766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DE Human PRO polypeptide #205.

DE Human PRO polypeptide #205.

PN US200199059-Al.

PD 23-OCT-2003.

PA (GETH ) GENENTECH INC.

Querry Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2002203433-A1.
                                                                                                                                                                                                                                                                                                                                                                                                    Human PRO polypeptide #205.
102003199033-A1.
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US2003199064-Al.
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30-000-203203429-A1.
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US2003199026-A1.
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Human PRO polypeptide #205.
US2003199032-A1.
                                                                                              ADE19516 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003199025-A1.
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23-OCT-2003.
(GETH ) GENENTECH INC.
76.0%;
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                             Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 164
                                                       Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                    Query Match 76.0%; Score 174; DB 7; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE33455 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE34007 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
TETY MATCh 76.0%; Score 174; DB 7; LK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76.0%; Score 174; DB 7; L 100.0%; Pred. No. 1.2e-158;
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A (GETH) GENENTECH INC.
Ouery Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-OCT-2003.
(GETH) GENENTECH INC.
Query Match 176.0%; Score 174; DB 7; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003194781-A1.
                                                                                                                  ADD72264 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GETH ) GENENTECH INC.
76.0%; Score 174;
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US2003199053-Al.
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US2003194772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE17860 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD80059 standard; protein; 229 AA. Human PRO polypeptide #205. US2003207417:A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD93096 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194768-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                        GENENTECH INC.
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Best Local Similarity
RESULT 168
                    30-OCT-2003
(GETH ) GEN
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Query Match 76.0%; Score 174; DB 7; Length. 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 182
ID ADE40923 standard; protein; 229 AA.
     76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                  76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 184
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   76.0%; Score 174; DB 7; Length 229; 100.0%; Pred. No. 1.2e-158;
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76.0%; Score 174; DB 7; Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207355-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG23201 standard, protein, 229 AA.
Novel human secreted and transmembrane protein PRO733.
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76.0%; Score 174; DB 7; Le

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(ATY DB 7; Le
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADF47287 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195333-A1.
                                                         ADD80¢11 standard; protein; 229 AA. Human PRO polypeptide #205. US2003207418-A1. G6-NOV-2003. (GETH ) GENENTECH INC.
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US2003199031-Al.
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US2003199034-Al.
                                                                                                                                                                                                       ADD89639 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003199028-Al.
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Human NOVX4c protein.
WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE92851 standard; protein; 229 AA Human PRO polypeptide #205.
US2003194777-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(CURA-) CURAGEN CORP.
Query Match
Best Local Similarity
RESULT 180
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RESULT 181
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1D ADP97536 standard; protein; 229 AA.

1D Human PRO polypeptide #205.

PN US2003207370-A1.

PD 06-NOV-2003

Query Match

RESULT 190

RESULT 190
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(GETH) GENENTECH INC.
76.0%; Score 174; DB 7; Length 229;
t Local Similarity 100.0%; Pred. No. 1.2e-158;
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Query Match 76.0%; Score 174; DB 7; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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ry Match 76.0%; Score 174; DB 7; Length 229;
t Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733
US2003207379-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM55340 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207381-A1.
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(GETH ) GENENTECH INC.
rry Match 76.0%; Score 174; DB 7; Le
rrann Similarity 100.0%; Pred. No. 1.2e-158;
Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 189
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(GETH) GENENTECH INC.
ery Match 76.0%; Score 174; DB 7; Lest Local Similarity 100.0%; Pred. No. 1.2e-158;
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Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003216561-A1.
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Human secreted/transmembrane protein, PRO733.
US2003206915-A1.
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US2003077700-A1.
                                                                                                                                                                                                                                                ADG80600 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207373-A1.
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Human PRO polypeptide #205.
US2003207372-A1.
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(GETH ) GENENTECH INC.
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ADC81147 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003092115-A1.
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(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 7; Length 229;
(ery Match 74milarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
Pery Match
Tonal Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                  ADI63559 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207387-A1.
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Novel human secreted and transmembrane protein PRO733.
US203087356-A1.
                    ADI65060 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADN16618 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADN14885 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003087357-A1.
                                                                                                                                                                                                                                                                                                                                                                                 ADH81973 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH81421 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207377-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM82590 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087355-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADNI5989 standard, protein, 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003087353-A1.
                                                                                                                                                                                                                                                                         06-NOV-zous.
(GETH) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 7; Le
2ry Match 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                   Best Local Similarity RESULT 199
                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 200
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RESULT 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                               US2003207388-A1.
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15-MAY-2003.
(GETH ) GENENTECH INC.
Query March
76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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76.0%; Score 174; DB 8; Length 229; t Local Similarity 100.0%; Pred. No. 1.2e-158;
PD 15-MAY-2003.
PA (GE-MAY-2003).
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 208
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
Query Match
Dest Local Similarity 100.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GFTH) GENENTECH INC.
176.0%; Score 174; DB 8; Lust Match
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003104536-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PRO polypeptide #205.
102003203440-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE75811 standard; protein; 229 AA. Human PRO polypeptide #205. US2003211571-A1.
                                                                                                                           ADD76595 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003100087-A1.
                                                                                                                                                                                                                                                                                                         ADD87959 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003092113-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE23387 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003092108-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE23939 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003092110-A1.
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US2003092111-A1.
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30-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; SCC
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(GETH ) GENENTECH INC.
76.0%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 209
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vuery Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 223
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PA (GETH.) GENENTECH INC.

Query Match 76.0%; Score 174; DB 8; Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 228
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(GETH) GENENTECH INC.
ery Match
et Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
2ry Match 76.0%; Score 174; DB 8; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.

Query Match 76.0%; Score 174; DB 8; Lo

Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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PA (CETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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ry Match

t Local Similarity 100.0%; Pred. No. 1.2e-158;
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( GETH ) GENENTECH INC.

76.0%; Score 174; DB 8; L

PEST Local Similarity 100.0%; Pred. No. 1.2e-158;
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A (GETH) GENENTECH INC.
76.0%; SCOTE.174; DB 8; L.
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
    ADF40214 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003198994-A1.
                                                                                                                                                                     ADF46010 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195148-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF24406 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003204055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADF40838 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199021-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF23782 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003203402-A1.
                                                                                                                                                                                                                                                                                                                                         ADE94741 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199027-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE91152 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE93403 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199060-A1.
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Human PRO polypeptide #205.
US2003199052-A1.
                                                                  23-OCT-2003.
(GETH ) GENENTECH INC.
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                   PD 30-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                         Length 229;
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                                                                                                                                                                                      PD 23-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; LA Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 218
                                                                                                                                                                                                                                                                                                                                                          PD 16-OCT-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 219
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(GETH ) GENENTECH INC.

Query Match
76.0%; Score 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003130181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADF61522 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003195345-A1.
                                                                                                                                                                                                                                                                                                  Human PRO polypeptide #205.
1022003194794-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE88721 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003199054-A1.
                                                                                                                                              ADE89273 standard; protein; 229 AA Human PRO polypeptide #205. US2003199062-A1.
Human PRO polypeptide #205.
US2003203439-A1.
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BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-OCT-2003.
(GETH ) GENENTECH INC.
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PAONI N F.
ROY M A.
SHELTON D L.
STEWART T A.
TUMAS D.
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DESNOYERS L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILVAROFF E
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KLJAVIN I J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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Best Local Similarity
RESULT 222
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NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EATON D L
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GERBER H.
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06-NOV-2003
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                                                                                                                Query Match
                             PD 16-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   76.0%; Score 174; DB 8; Length 229; 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003199051-A1.
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(GETH ) GENENTECH INC.
(ery Match 76.0%; Score 174; DB 8; Lk
(ery Match 100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 233
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(GETH ) GENENTECH INC.
16.0%; Score 174; DB 8; L.
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100.0%; Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003199435-A1.
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Human secreted/transmembrane protein, PRO733.
US2003211091-A1.
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Human secreted/transmembrane protein, PRO733.
US2003211092-A1.
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Human secreted/transmembrane protein, PRO733.
Human secreted/transmembrane protein, PRO733.
US2003194780-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF27232 standard; protein; 229 AA..
Human secreted/transmembrane protein, PRO733.
US2003199436-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADF27868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2003199437-A1.
                                                                                                                                                                                                                                               ADF34984 standard; protein; 229 AA. Human PRO polypeptide #205.
US2003199029-A1.
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US2003199063-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local
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(GETH ) GENENTECH INC.

ry Match 76.0%; Score 174; DB 8; Length 229; Local Similarity 100.0%; Pred. No. 1.2e-158;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 229;
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(GETH) GENENTECH INC.
76.0%; Score 174; DB 8; Length 229;
t Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                          Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207426-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE91747 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003199058-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207360-A1.
US2003199674-A1.
23-OCT-2003.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; Le
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ry Match
t Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                           DE Human secreted/transmembrane protein, PRO733.
PN US2003194410-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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A (GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; L.
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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4 (GETH ) GENENTECH INC.

Query Match

76.0%; Score 174; DB 8;

Best Local Similarity 100.0%; Pred. No. 1.2e-156
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Human secreted/transmembrane protein, PRO733
US2003195344-A1.
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0S2003207376-A1.
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Human PRO polypeptide #205.
US2003207352-A1.
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Human PRO polypeptide #205.
US2003207422-A1.
06-NOV-2003.
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Human PRO polypeptide #205.
US2003208055-A1.
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06-NOV-2003.
(GETH ) GENENTECH INC.
76.0%; SCO
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(GETH ) GENENTECH INC.
                                                                                                          Best Local Similarity RESULT 241
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DB 8; Length 229;

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OG-NOV-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Length 229;
ery Match 100.0%; Pred. No. 1.2e-158;
                  (GETH ) GENENTECH INC.
1:17 Match
1:1 Ancal Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
Query Match
76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
Pry Match 76.0%; Score 174; DB 8; Length 229;
st Local Similarity 100.0%; Pred. No. 1.2e-158;
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B Novel human secreted and transmembrane protein PRO733.

PN US2003207356-A1.

PD 06-NOV-2003.

PA (GETH) GENENTECH INC.

Query Match 76.0%; Score 174; DB 8; Length Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733
US2003207389-A1.
GG-NOV-2003
(GETH ) GENENTECH INC.
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US2003207427-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207428-AI.
(GE-NOV-2007)
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG55287 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003194778-A1.
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 264
                                                                                                      ADG06169 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207374-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                  ADG04042 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003207423-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                            Query Match
Best Local Similarity
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Best Local Similarity
RESULT 266
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PD 06-NOV-2003.
PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 250
                                                                                                                  ADC03490 standard; protein; 229 AA.

Human PRO polypeptide #205.
US2003207351-A1.
06-NOV-2003.
(GETH ) GENENTECH INC.
Fry Match
T6.0%; Score 174; DB 8; Length 229;
St Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
ery Match
76.0%; Score 174; DB 8; Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207424-A1.
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 253
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GG-NQC-NQC-2003.

(GETH ) GENENTECH INC.

76.0%; Score 174; DB 8; L.

100.0%; Pred. No. 1.2e-158;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 254
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 255
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US2003207353-A1.
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US2003207375-A1.
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US2003207425-A1.
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Human PRO polypeptide #205.
US2003207357-A1.
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US2003219885-A1.
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Human PRO polypeptide #205.
US2003207371-A1.
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2003.
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2ry Match
Best Local Similarity 1v.
RESULT 256
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PN US20032.
PD 06-"
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 252
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Best Local Similarity
RESULT 251
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BERRE

Length 229;

Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 277

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Query Match 76.0%; Score 174; DB 8; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 272
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  Length 229;
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Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003207415-A1.
                                                                                                                                                                                                                                         ADG57495 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207362-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG70965 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207420-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADGS8047 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207363-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207421-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                    ADGS6943 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2003207364-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207365-A1.
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Novel human secreted and transmembrane protein PRO733
US2003207368-A1.
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                                                                                                                                                                                                                                                                                                                                                      76.0%; Score 174; DB 8; Le
100.0%; Pred. No. 1.2e-158;
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  76.0%; Score 174; DB 8; L 100.0%; Pred. No. 1.2e-158;
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                                                                 ADG82256 standard; protein; 229 AA
                                                                                                                                                                            76.0%; S
100.0%;
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                                                                                      Human PRO polypeptide #205.
US2003207358-A1.
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Query Match
Best Local Similarity
RESULT 268
1D ADG82256 standard, pro
DE Human PRO polypeptide
PN US2003207358-A1.
PD 06-NOV-2003.
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RESULT 270
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vuery Match 76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 281
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ry Match 76.0%; Score 174; DB 8; Length 229; t Local Similarity 100.0%; Pred. No. 1.2e-158;
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Local Similarity 100.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
US2003207419-A1.
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Novel human secreted and transmembrane protein PRO733.
US200207416-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG52455 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207414-A1.
                                                                    PA (GETH) GENENTECH INC.
Query Match 76.0%; Score 174; DB 8; Lv Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 278
                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 174; DB 8; Louised. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003215905-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein, PRO733.012206305-A1.
ADG50620 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003207803-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG51868 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733
US2003215908-A1.
                                                                                                                                                                                                                                                                                                                       ADH30666 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003077723-A1.
24-APR-2003.
                                                                                                                                                                     ADG81704 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2003207805-A1.
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(GETH ) GENENTECH INC.
76.0%;
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
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(GETH ) GENENTECH INC.
(GETY Match 76.0%; Score 174; DB 8; Length 229;
ery Match 76.0%; Pred. No. 1.2e-158;
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Novel human secreted and transmembrane protein PRO733.
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US2003207366-A1.
                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PR0733 US2003207378-A1.
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Novel human secreted and transmembrane protein PRO733.
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NOG11 human secreted and transmembrane protein PRO733.
US2003207367-A1.
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20-NOV-2003.
(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; La
ery Match 100.0%; Pred. No. 1.2e-158;
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Pred. No. 1.2e-158;
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100.0%; Pred. No. 1.2e-158;
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Human secreted/transmembrane protein, PRO733.
US2003216560-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADGS1244 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004005312-A1.
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30-JAN-2003.
(GETH) GENENTECH INC.
GEV MALCh
AGENTITY 100.0%; Pred. No.
ADG81152 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003194793-A1.
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                                                                                                                                      protein; 229 AA.
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06-NOV-2003.
(GETH ) GENENTECH INC.
""= Ch 76.0%; S.
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2.4 ) GENENTECH IN
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 289
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RESULT 294
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                                                                                                                                    ADG56391 standard;
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ADH25669 standard; protein; 229 AA.
Human neurotrimin homologue related protein sequence SEQ ID NO:447.
                                                                                                  DB 8; Length 229;
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Novel human secreted and transmembrane protein PRO733.
US2003207382-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207383-A1.
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Novel human secreted and transmembrane protein PRO733.
US2003207349-A1.
06-NOV-2003.
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Query Match 76.0%; Score 174; DB 8; L4

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 298
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(GETH ) GENENTECH INC.
ery Match 76.0%; Score 174; DB 8; Leery Match 76.0%; Pred. No. 1.2e-158;
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76.0%; Score 174; DB 8; Lv
al Similarity 100.0%; Pred. No. 1.2e-158;
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( GETH ) GENENTECH INC.

76.0%; Score 174; DB 8; Lv

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Best Local Similarity 100.0%; Pred. No. 1.2e-158;
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(GETH) GENENTECH INC.
QUERY MAICH 76.0%; SCOTE 174; DB 8; L
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 8;
Human secreted/transmembrane protein, PRO733.
US2004005657-A1.
08-JAN-2004.
                                                                                                                                                               ADG62644 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004006219-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 174;
                                                                                                                                                                                                                                                                                                                                             ADI81199 standard; protein; 229 AA. Human PRO polypeptide #205. US2003207361-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             )6-NOV-ZUUS
(GETH ) GENENTECH INC.
76.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    76.0%;
                                                                                                                                                                                                                              08-JAN-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG09942 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG09290 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EP1386931-A1.
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A (GETH) GENENTECH INC.
Query Match
76.0%; Score 174; DB 8; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 76.0%; Score 174; DB 8; Length 229; Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    76.0%; Score 174; DB 9; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                           / Match 76.0%; Score 174; DB 8; Length 229; Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel bronchial cancer-associated human protein SeqID899.
DE10316701-A1.
                                                                                                                                            ADS323346 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733
US2004203125-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11.NOV-2004.
(GETH) GENENTECH INC.
PY MATCH
T. Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         / Match 76.0%; Score 174; DB 8; L. Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide #205.
US2004214269-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADU50402 standard; protein; 229 AA.
PRO733, SEQ ID 447.
US2004223964-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADU06673 standard; protein; 229 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HINZ/) HINZMANN B.
(HERM/) HERMANN K.
(CAST/) HEIDEN CASTANOS-VELEZ E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-0CT-2004
(GETH ) GENENTECH INC. 76.0%;
                                                                                                                                                                                                                    USACOT-2004.
14-OCT-2004.
(GETH ) GENENTECH INC.
76.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRO733 protein, SEQ ID 447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASHKENAZI A J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TUMAS D.
WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ROY M A.
SHELTON D L.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EATON D L.
FERRARA N.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 319
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PAONI N F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DESNOYERS 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KUO S S.
NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADW49681 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GODDARD A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GURNEY A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HILLAN K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS2005014226-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .
⊗
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     06-NOV-2003
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                                                                                                                                                                                                                                                                                              Query Match
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(WOOD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SHEL/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PANJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 229;
                                                                                      76.0%; Score 174; DB 8; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                            Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 229;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AD196394 standard; protein; 229 AA.
Novel human secreted and transmembrane protein PRO733.
US2003207354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-APR-2004.
(GETH ) GENENTECH INC.
76.0%; Score 174; DB 8; LE.
100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USZUCTO...
USZUCTO...
(GETH ) GENENTECH INC.
APTV MALCh ...
APTV M
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PA (GETH) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 100.0%; Score 1/4; DB 8; L RESULT 311
                                                                                                                                                                                                                                                                                                    Query Match 76.0%; Score 174; DB 8; L Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PD 11-MAR-2004.

PA (GETH ) GENENTECH INC.

Query March

Best Local Similarity 100.0%; Pred. No. 1.2e-158;

RESULT 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 76.0%; Score 174; DB 8; L. Best Local Similarity 100.0%; Pred. No. 1.2e-158; RESULT 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-2004.
(GETH ) GENENTECH INC.
76.0%; - Score 174; DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADMI7446 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004048332-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL07280 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2004063921-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76.0%; Score 174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human PRO polypeptide #205.
22-app.
                                                                                                                                                           Human PRO polypeptide #205.
US2004038336-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          ADJ65638 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2004038335-A1.
26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADM27774 standard, protein, 229 AA.
Human PRO polypeptide #205.
US2004048333-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AD195842 standard; protein; 229 AA.
Human PRO polypeptide #205.
US2003077659-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM42498 standard; protein; 229 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.08;
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US2004058424-Al.
US2004039164-A1.
26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                   26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GETH ) GENENTECH INC
                                                                                                           Best Local Similarity RESULT 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-APR-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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Length 229;

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06-APR-2006.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           13-APR-2006. (GETH ) GENENTECH INC.
                                                                               STEWART T A.
TUMAS D.
WATANABE C K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GERRITSEN M E.
GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STEWART T A.
TUMAS D.
WATANABE C K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DEFORGE L.
DESNOYERS L.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GODOWSKI P J
                                                                                                                                                             Query Match
Best Local Similarity
RESULT 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 329
                                     GURNEY A L. SHERWOOD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SHERWOOD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GURNEY A L.
        GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BAKE/) BAKER K P.
                                                                    SMITH V.
                                                                                                                                                ZHANG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SMITH V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZHANG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-FEB-2006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A (WOOD/) WC
A (ZHAN/) ZH
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                 (MOOD/)
(ZHAN/)
                                                                    (SMIT/)
(STEW/)
                                                                                                  (TUMA/)
(WATA/)
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(TUMA/)
(WATA/)
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(DESN/)
(FILV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SHER/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (/LIWS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AEB14127 standard; protein; 229 AA.
Cancer cell diagnosis method-related human protein - SEQ ID 410.
US2005153396-A1.
                                   PD 24-FEB-2005.
PA (GERENTECH INC.
Query Match 76.0%; Score 174; DB 9; Length 229;
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
RESULT 320
                                                                                                                                                                                                            Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  76.0%; Score 174; DB 9; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                           PD 07-APR-2005.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          76.0%; Score 174; DB 9; L 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                      ADZ52342 standard; protein; 229 AA.
Human secreted/transmembrane protein, PRO733.
US2005084935-A1.
                                                                                                                            ADZ03381 standard; protein; 229 AA.
Human secreted/transmembrane PRO733 protein.
US2005074837-A1.
ADY14737 standard; protein; 229 AA.
PRO polypeptide SEQ ID NO 543.
WO2005016962-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AEA99614 standard; protein; 229 AA.
Human PRO733 protein SEQ ID NO:2.
US2005118635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JO/)
JM/)
JM/) GD.,
SMIT/) SMITH
(WOOD/) WOOD W I.,
Lery Match
JSULT 323
LD AEB14127 standard;
DE Cancer cell diagr
PN US2005153396-A1
PD 14-JUL-2005.
PA (BAKE/) BF
PA (BREK/) BF
PA (BREK/)
PA (FII')
PA (FII')
PA (FII')
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GODD/) GODDARD A. (GODO/) GODOWSKI P J. (GURN-) GURNEY A L. (SMIT/) SMITH V. (WODD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                     ASHKENAZI A J.
BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TUMAS D.
WILLIAMS P.M.
WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ROY M A..
SHELTON D L.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                  BOTSTEIN D.
DESNOYERS L.
EATON D L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODOWSKI P J
GRIMALDI J C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                   FILVAROFF E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KLJAVIN I J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PAONI N F.
                                                                                                                                                                                                                                                                                                                                                                                                   PERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODDARD A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GURNEY A HILLAN K
                                                                                                                                                                                                                                                                                                                                                                                                                                                               GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KUO S S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SAO W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-JUN-2005
                                                                                                                                                                                                                                                                                                                                     BAKE/
                                                                                                                                                                                                                                                                                                                                                                                                   FERR/
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PA (GODO) GODOWSKI P J.

PA (GIDNA) GODOWSKI P J.

PA (SHRIA) SHEWAR I P J.

PA (SHRIA) SHEWAR I P A.

PA (STRAM) STRAWE I P A.

PA (GTTM) CHANNER I P A.

PA (GTTM) PILVAOFF E.

PA (GTTM) PILVAOFF E.

PA (GTTM) CHANNER I P A.

PA (GTTM) SHARE I P A.

PA (GTTM) SHARE I P A.

PA (GTTM) SHARE I P A.

PA (GTTM) CHANNER I P A.

PA (G
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Extended human secreted protein sequence, SEQ ID NO. 149.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                         Query Match
                                                                          Ouery Match
                                                                                            мыл7411 standard; protein; 229, АА.
Human cancer treatment biomarker amino acid sequence - SEQ ID 298.
US2006121511-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADL22684 standard; protein; 141 AA.
Human disease detection and treatment (MDDT) protein - SEQ ID 133.
WO2003062379-A2.
                                                                                                                                                                                                                                                                                                                                                  21-JUN-2006.
(GETH) GENENTECH INC.
ery Match
-- ' . A similarity 100.0%; Pred. No. 1.2e-158;
                                                     76.0%; Score 174; DB 10; Length 229; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                          Length 229,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 56.3%; Score 129; DB 7; Length 229; 100.0%; Pred. No. 2.3e-115;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 8; Length 229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68.6%; Score 157; DB 7; Length 162; 100.0%; Pred. No. 2e-142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 133; DB 7; Length 141;
Pred. No. 2.2e-119;
                                                                                                                                                                                                                                                                                      69.9%; Score 160; DB 2; Length 229; 100.0%; Pred. No. 3.6e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 193;
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY35962 standard; protein; 229 AA.
Extended human secreted protein sequence, SEQ ID NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68.6%; Score 157; DB 7; Le 100.0%; Pred. No. 2.4e-142;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 3.6e-145;
                                                                                                                                                                                                                                          76.0%; Score 174; DB 10; 100.0%; Pred. No. 1.2e-158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               69.9%; Score 160; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE95518 standard; protein; 162 AA.
Human NOVX4b protein.
WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG14988 standard; protein; 193 AA.
Human SECP-21 protein.
WO2003087300-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE95516 standard, protein, 229 AA.
Human NOVX4a protein.
WO2003050245-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADP19270 standard; protein; 229 AA. Human secreted polypeptide #121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY35900 standard; protein; 162 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INCY-) INCYTE GENOMICS INC. ry Match 58.1%; t Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INCY-) INCYTE CORP.
                                                                                                                                                                 08-JUN-2006.
(LEEH/) LEE H.
(SHAW/) SHAW P.
(CLAR/) CLARK E A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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RESULT 337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 335
                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 331
                                                                         Best Local Similarity
(LEEH/) LEE H.
(SHAW/) SHAW P.
(CLAR/) CLARK E A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GEST ) GENSET SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-JUN-1999.
(GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9931236-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-JUN-200
                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouerv Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 336
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AB023233 standard, protein; 169 AA.
Human breast tumour associated protein 47-like polypeptide NOV1
US2003027158-A1.
                                                           54.6%; Score 125; DB 2; Length 162 100.0%; Pred. No. 1.2e-111;
                                                                                                                                                                                                                                 Length 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   34.9%; Score 80; DB 8; Length 169; 100.0%; Pred. No. 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 8; Length 169; 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 6; Length 169;
2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        34.9%; Score 80; DB 7; Length 169; 100.0%; Pred. No. 2.4e-68;
                                                                                                                                                                                                                                                                                                                                                                                                    41.9%; Score 96; DB 2; Length 120; 100.0%; Pred. No. 7.2e-84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 4; Length 169;
2.4e-68;
                                                                                                                                                                                                                                                                                          AAY35901 standard; protein; 120 AA.
Extended human secreted protein sequence, SEQ ID NO. 150.
WO9931236-A2.
                                                                                                                                                                                                                               54.6%; Score 125; DB 8; I 100.0%; Pred. No. 1.2e-111;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 41.9%; Score 96;
Best Local Similarity 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 34.9%; Score 80;
Best Local Similarity 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34.9%; Score 80; 100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 34.9%; Score 80; Best Local Similarity 100.0%; Pred. No.
                                                                                                       Human secreted polypeptide #59. US2004110939-A1. 10-UN-2004. GEST ) GENSET SA. ry Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP19209 standard; protein; 120 AA.
Human secreted polypeptide #60.
US2004110939-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB61129 standard; protein; 169 AA.
Human NOV1 protein.
WO200075321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADMS6365 standard; protein; 169 AA. Human cell adhesion molecule NOV1. US2003082554-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI19766 standard; protein; 169 AA.
Human NOV1 protein.
US2004002134-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF66729 standard; protein; 169 AA.
Novel human protein NOV1.
US2003199103-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADO60239 standard; protein; 169 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-DEC-2000.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-FEB-2003.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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Best Local Similarity
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                                                                              Local Similarity
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                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JUN-2004.
(GEST ) GENSET SA.
                                                                                                                                                                                                                                                                                                                                                        24-JUN-1999.
(GEST ) GENSET.
                   24-JUN-1999.
(GEST ) GENSET.
WO9931236-A2.
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US2003134430-A1.

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PD 17-UL-2003.
PD 17-UL-2003.
PD 17-UL-2003.
PD 17-UL-2003.
PD 27-UL-2003.
PA (CURA.) Added the Curacar Corp.
PA (CURA.) Added the Curacar Curacar No. 2.46-68;
PRESULT 348 | 100.04; Pred. No. 2.46-68;
PA (GENY ) GENETICS INST INC.
OMENY MAION GENETICS INST INC.
DAMBIO48 standard; protein; 133 AA.
DE Human adult blood protein fragment AIG-11.
PN WGO20037630-AI.
PN WGO20037630-AI.
PN WGO20037630-AI.
PN WGO20037630-AI.
PN WGO20037630-AI.
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PN WGO30037630-AI.
PN WGO300530-AI.
PN WGO300141-AIR
PN WGO30014-AIR
PN WGO300014-AIR
PN WGO30014-AIR
PN WGO300014-AIR
PN WGO300014-AIR
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